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FBA object

The FBA object is a collection of nodes in a scene graph which are animated by the FBA (Face and Body Animation) object bitstream. The FBA object is controlled by two separate bitstreams. The first bitstream, called BIFS, contains instances of Body Definition Parameters (BDPs) in addition to Facial Definition Parameters (FDPs) , and the second bitstream, FBA bitstream, contains Body Animation Parameters (BAPs) together with Facial Animation Parameters (FAPs).

A 3D (or 2D) *face object* is a representation of the human face that is structured for portraying the visual manifestations of speech and facial expressions adequate to achieve visual speech intelligibility and the recognition of the mood of the speaker. A face object is animated by a stream of *face animation parameters (FAP)* encoded for low-bandwidth transmission in broadcast (one-to-many) or dedicated interactive (point-to-point) communications. The FAPs manipulate key feature control points in a mesh model of the face to produce animated visemes for the mouth (lips, tongue, teeth), as well as animation of the head and facial features like the eyes. FAPs are quantized with careful consideration for the limited movements of facial features, and then prediction errors are calculated and coded arithmetically. The remote manipulation of a face model in a terminal with FAPs can accomplish lifelike visual scenes of the speaker in real-time without sending pictorial or video details of face imagery every frame.

A simple streaming connection can be made to a decoding terminal that animates a default face model. A more complex session can initialize a custom face in a more capable terminal by downloading *face definition parameters (FDP)* from the encoder. Thus specific background images, facial textures, and head geometry can be portrayed. The composition of specific backgrounds, face 2D/3D meshes, texture attribution of the mesh, etc. is described in ISO/IEC 14496-1. The FAP stream for a given user can be generated at the user's terminal from video/audio, or from text-to-speech. FAPs can be encoded at bitrates up to 2-3kbit/s at necessary speech rates. Optional temporal DCT coding provides further compression efficiency in exchange for delay. Using the facilities of ISO/IEC 14496-1, a composition of the animated face model and synchronized, coded speech audio (low-bitrate speech coder or text-to-speech) can provide an integrated low-bandwidth audio/visual speaker for broadcast applications or interactive conversation. Limited scalability is supported. Face animation achieves its efficiency by employing very concise motion animation controls in the channel, while relying on a suitably equipped terminal for rendering of moving 2D/3D faces with non-normative models held in local memory. Models stored and updated for rendering in the terminal can be simple or complex. To support speech intelligibility, the normative specification of FAPs intends for their selective or complete use as signaled by the encoder. A masking scheme provides for selective transmission of FAPs according to what parts of the face are naturally active from moment to moment. A further control in the FAP stream allows face animation to be suspended while leaving face features in the terminal in a defined quiescent state for higher overall efficiency during multi-point connections.

A body model is a representation of a virtual human or human-like character that allows portraying body movements adequate to achieve nonverbal communication and general actions. A body model is animated by a stream of *body animation parameters* (BAP) encoded for low-bitrate transmission in broadcast and dedicated interactive communications. The BAPs manipulate independent degrees of freedom in the skeleton model of the body to produce animation of the body parts. The BAPs are quantized considering the joint limitations, and prediction errors are calculated and coded arithmetically. Similar to the face, the remote manipulation of a body model in a terminal with BAPs can accomplish lifelike visual scenes of the body in real-time without sending pictorial and video details of the body every frame.

The BAPs, if correctly interpreted, will produce reasonably similar high level results in terms of body posture and animation on different body models, also without the need to initialize or calibrate the model.The BDP set defines the set of parameters to transform the default body to a customized body optionally with its body surface, body dimensions, and texture.

The *body definition parameters (BDP)* allow the encoder to replace the local model of a more capable terminal. BDP parameters include body geometry, calibration of body parts, degrees of freedom, and optionally deformation information.

The Face Animation specification is defined in ISO/IEC 14496-1 and this part of ISO/IEC 14496. This clause is intended to facilitate finding various parts of specification. As a rule of thumb, FAP specification is found in the part 2, and FDP specification in the part 1. However, this is not a strict rule. For an overview of FAPs and their interpretation, read subclauses "6.1.5.2 Facial animation parameter set", "6.1.5.3 Facial animation parameter units", "6.1.5.4 Description of a neutral face" as well as the Table C-1. The viseme parameter is documented in subclause "7.12.3 Decoding of the viseme parameter fap 1" and the Table C-5 in annex C. The expression parameter is documented in subclause "7.12.4 Decoding of the expression parameter fap 2" and the Table C-3 . FAP bitstream syntax is found in subclauses "6.2.10 Face Object", semantics in "6.3.10 Face Object", and subclause "7.12 Face object decoding" explains in more detail the FAP decoding process. FAP masking and interpolation is explained in subclauses "6.3.11.1 Face Object Plane", "7.12.1.1 Decoding of faps", "7.12.5 Fap masking" . The FIT interpolation scheme is documented in subclause "7.2.5.3.2.4 FIT" of ISO/IEC 14496-1. The FDPs and their interpretation are documented in subclause "7.2.5.3.2.6 FDP" of ISO/IEC 14496-1. In particular, the FDP feature points are documented in Figure C-1.

FBA object

Conceptually the FBA object consists of a collection of nodes in a scene graph which are animated by the facial object bitstream. The shape, texture and expressions of the face are generally controlled by the bitstream containing instances of Facial Definition Parameter (FDP) sets and/or Facial Animation Parameter (FAP) sets. Upon construction, the FBA object contains a generic face with a neutral expression. This face can already be rendered. It is also immediately capable of receiving the FAPs from the bitstream, which will produce animation of the face: expressions, speech etc. If FDPs are received, they are used to transform the generic face into a particular face determined by its shape and (optionally) texture. Optionally, a complete face model can be downloaded via the FDP set as a scene graph for insertion in the face node.

The FDP and FAP sets are designed to allow the definition of a facial shape and texture, as well as animation of faces reproducing expressions, emotions and speech pronunciation. The FAPs, if correctly interpreted, will produce reasonably similar high level results in terms of expression and speech pronunciation on different facial models, without the need to initialize or calibrate the model. The FDPs allow the definition of a precise facial shape and texture in the setup phase. If the FDPs are used in the setup phase, it is also possible to produce more precisely the movements of particular facial features. Using a phoneme/bookmark to FAP conversion it is possible to control facial models accepting FAPs via TTS systems. The translation from phonemes to FAPs is not standardized. It is assumed that every decoder has a default face model with default parameters. Therefore, the setup stage is not necessary to create face animation. The setup stage is used to customize the face at the decoder.

Upon construction, the Body object contains a generic virtual human or human-like body with the default posture. This body can already be rendered. It is also immediately capable of receiving the BAPs from the bitstream, which will produce animation of the body. If BDPs are received, they are used to transform the decoder's generic body into a particular body determined by the parameter contents. Any component can be null. A null component is replaced by the corresponding default component when the body is rendered. Similar to the face, the BAPs can be transmitted also without first downloading BDPs, in which case the decoder animates its local model.

No assumption is made and no limitation is imposed on the range of defined mobilities for humanoid animation. In other words the human body model should be capable of supporting various applications, from realistic simulation of human motions to network games using simple human-like models.

Structure of the face and body object bitstream

A face and body object is formed by a temporal sequence of face and body object planes. This is depicted as follows in Figure 6-9.

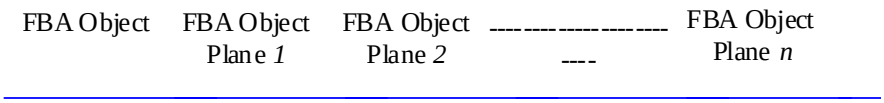
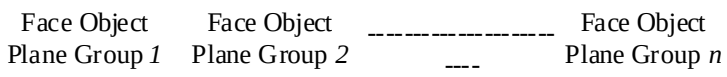


Figure 6-9 -- Structure of the FBA object bitstream

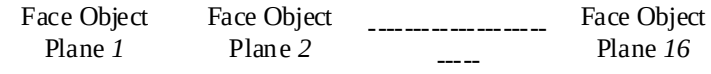
An FBA object represents a node in an ISO/IEC 14496 scene graph. An ISO/IEC 14496 scene is understood as a composition of Audio-Visual objects according to some spatial and temporal relationships. The scene graph is the hierarchical representation of the ISO/IEC 14496 scene structure (see ISO/IEC 14496-1).

Alternatively, an FBA object can be formed by a temporal sequence of FBA object plane groups (called segments for simplicity), where each FBA object plane group itself is composed of a temporal sequence of 16 FBA object planes, as depicted in the following:

FBA **object:**



FBA **object plane group:**



When the alternative FBA object bitstream structure is employed, the bitstream is decoded by DCT-based FBA object decoding as described in subclause 7.12.2. Otherwise, the bitstream is decoded by the frame-based FBA object decoding. Refer to Table C-1 for a specification of default minimum and maximum values for each FAP

Facial animation parameter set

The FAPs are based on the study of minimal facial actions and are closely related to muscle actions. They represent a complete set of basic facial actions, and therefore allow the representation of most natural facial expressions. Exaggerated values permit the definition of actions that are normally not possible for humans, but could be desirable for cartoon-like characters.
The FAP set contains two high level parameters visemes and expressions. A viseme is a visual correlate to a phoneme. The viseme parameter allows viseme rendering (without having to express them in terms of other parameters) and enhances the result of other parameters, insuring the correct rendering of visemes. Only static visemes which are clearly distinguished are included in the standard set. Additional visemes may be added in future extensions of the standard. Similarly, the expression parameter allows definition of high level facial expressions. The facial expression parameter values are defined by textual descriptions. To facilitate facial animation, FAPs that can be used together to represent natural expression are grouped together in FAP groups, and can be indirectly addressed by using an expression parameter. The expression parameter allows for a very efficient means of animating faces. In annex C, a list of the FAPs is given, together with the FAP grouping, and the definitions of the facial expressions.

Facial animation parameter units

All the parameters involving translational movement are expressed in terms of the *Facial Animation Parameter Units (FAPU).* These units are defined in order to allow interpretation of the FAPs on any facial model in a consistent way, producing reasonable results in terms of expression and speech pronunciation. They correspond to fractions of distances between some key facial features and are defined in terms of distances between feature points. The fractional units used are chosen to allow enough precision. annex C contains the list of the FAPs and the list of the FDP feature points. For each FAP the list contains the name, a short description, definition of the measurement units, whether the parameter is unidirectional (can have only positive values) or bi-directional, definition of the direction of movement for positive values, group number (for coding of selected groups), FDP subgroup number (annex C) and quantisation step size. FAPs act on FDP feature points in the indicated subgroups. The measurement units are shown in Table 6-1, where the notation 3.1.y represents the y coordinate of the feature point 3.1; also refer to Figure 6-10.

Table 6-1 -- Facial Animation Parameter Units

Description		FAPU Value
IRISD0 = 3.1.y – 3.3.y = 3.2.y – 3.4.y	Iris diameter (by definition it is equal to the distance between upper ad lower eyelid) in neutral face	IRISD = IRISD0 / 1024
ES0 = 3.5.x – 3.6.x	Eye separation	ES = ES0 / 1024
ENS0 = 3.5.y – 9.15.y	Eye - nose separation	ENS = ENS0 / 1024
MNS0 = 9.15.y – 2.2.y	Mouth - nose separation	MNS = MNS0 / 1024
MW0 = 8.3.x – 8.4.x	Mouth width	MW = MW0 / 1024
AU	Angle Unit	10^{-5} rad

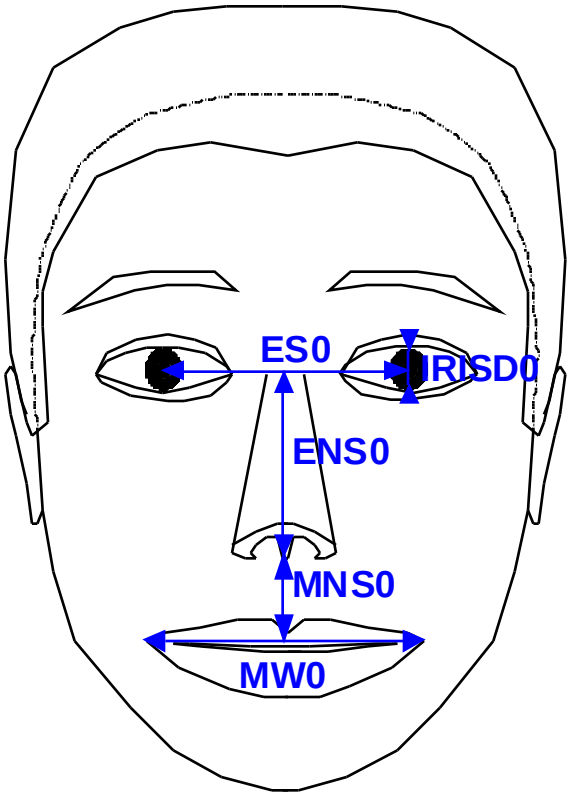


Figure 6-10 -- The Facial Animation Parameter Units

Description of a neutral face

At the beginning of a sequence, the face is supposed to be in a neutral position. Zero values of the FAPs correspond to a neutral face. All FAPs are expressed as displacements from the positions defined in the neutral face. The neutral face is defined as follows:

- the coordinate system is right-handed; head axes are parallel to the world axes
- gaze is in direction of Z axis
- all face muscles are relaxed
- eyelids are tangent to the iris
- the pupil is one third of IRISD0
- lips are in contact; the line of the lips is horizontal and at the same height of lip corners
- the mouth is closed and the upper teeth touch the lower ones
- the tongue is flat, horizontal with the tip of tongue touching the boundary between upper and lower teeth (feature point 6.1 touching 9.11 in annex C)

Facial definition parameter set

The FDPs are used to customize the proprietary face model of the decoder to a particular face or to download a face model along with the information about how to animate it. The definition and description of FDP fields is given in annex C. The FDPs are normally transmitted once per session, followed by a stream of compressed FAPs. However, if the decoder does not receive the FDPs, the use of FAPUs ensures that it can still interpret the FAP stream. This insures minimal operation in broadcast or teleconferencing applications. The FDP set is specified in BIFS syntax (see ISO/IEC 14496-1). The FDP node defines the face model to be used at the receiver. Two options are supported:

- calibration information is downloaded so that the proprietary face of the receiver can be configured using facial feature points and optionally a 3D mesh or texture.
- a face model is downloaded with the animation definition of the Facial Animation Parameters. This face model replace the proprietary face model in the receiver.

Body animation parameter set

BAP parameters comprise joint angles connecting different body parts. These include: toe, ankle, knee, hip, spine (C1-C7, T1-T12, L1-L5), shoulder, clavicle, elbow, wrist, and the hand fingers. The detailed joint list, with the rotation normals, are given in the following subclause. The rotation angles are assumed to be positive in the counterclockwise rotation direction with respect to the rotation normal. The rotation angles are defined as zero in the default posture, as defined below.

Note that the normals of rotation move with the body, and they are fixed with respect to the parent body part. That is to say, the axes of rotation are not aligned with the body or world coordinate system, but move with the body parts.

The hands are capable of performing complicated motions and are included in the body hierarchy. There are totally 29 degrees of freedom on each hand, assuming that the hand has a standard structure with five fingers.

The unit of rotations (BAPU) is defined as PI/10E-5 radians. The unit of translation BAPs (BAPs HumanoidRoot_tr_vertical, HumanoidRoot_tr_lateral, HumanoidRoot_tr_frontal) is defined in millimeters. The allowed range of BAPs is –10E5..10E5 BAPUs (-180..180 degrees).

Description of the default posture of the body

The default posture is defined by standing posture. This posture is defined as follows: the feet should point to the front direction, the two arms should be placed on the side of the body with the palm of the hands facing inward. This posture also implies that all BAPs have default values as 0.

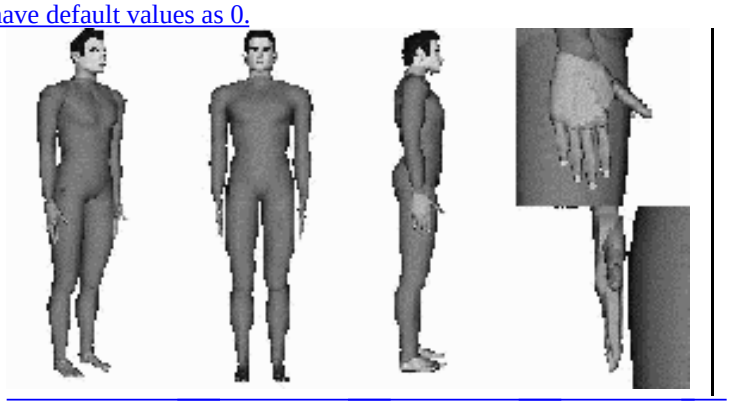


Figure V2 - 1

Visual bitstream syntax

Start codes

Start codes are specific bit patterns that do not otherwise occur in the video stream.

Each start code consists of a start code prefix followed by a start code value. The start code prefix is a string of twenty three bits with the value zero followed by a single bit with the value one. The start code prefix is thus the bit string '0000 0000 0000 0000 0000 0001'.

The start code value is an eight bit integer which identifies the type of start code. Many types of start code have just one start code value. However video_object_start_code and video_object_layer_start_code are represented by many start code values.

All start codes shall be byte aligned. This shall be achieved by first inserting a bit with the value zero and then, if necessary, inserting bits with the value one before the start code prefix such that the first bit of the start code prefix is the first (most significant) bit of a byte. For stuffing of 1 to 8 bits, the codewords are as follows in Table 6-2.

Table 6-2-- Stuffing codewords

Bits to be stuffed	Stuffing Codeword
1	0
2	01
3	011
4	0111
5	01111
6	011111

7	0111111
8	01111111

Table 6-3 defines the start code values for all start codes used in the visual bitstream.

Table 6-3 — Start code values

name	**start code value (hexadecimal)**
video_object_start_code	00 through 1F
video_object_layer_start_code	20 through 2F
reserved	30 through AF
visual_object_sequence__start_code	B0
visual_object_sequence_end_code	B1
user_data_start_code	B2
group_of_vop_start_code	B3
video_session_error_code	B4
visual_object_start_code	B5
vop_start_code	B6
reserved	B7-B9
fba_object_start_code	BA
fba_object_plane_start_code	BB
mesh_object_start_code	BC
mesh_object_plane_start_code	BD
still_texture_object_start_code	BE
texture_spatial_layer_start_code	BF
texture_snr_layer_start_code	C0
texture_tile_start_code	C1
texture_shape_layer_start_code	C2
reserved	C3-C5
System start codes (see note)	C6 through FF
NOTE System start codes are defined in ISO/IEC 14496-1	

The use of the start codes is defined in the following syntax description with the exception of the video_session_error_code. The video_session_error_code has been allocated for use by a media interface to indicate where uncorrectable errors have been detected.
This syntax for visual bitstreams defines two types of information:

1. Configuration information

   a. Global configuration information, referring to the whole group of visual objects that will be simultaneously decoded and composited by a decoder (VisualObjectSequence()).

   b. Object configuration information, referring to a single visual object (VO). This is associated with VisualObject().

   c. Object layer configuration information, referring to a single layer of a single visual object (VOL)

VisualObjectLayer()

2. Elementary stream data, containing the data for a single layer of a visual object.

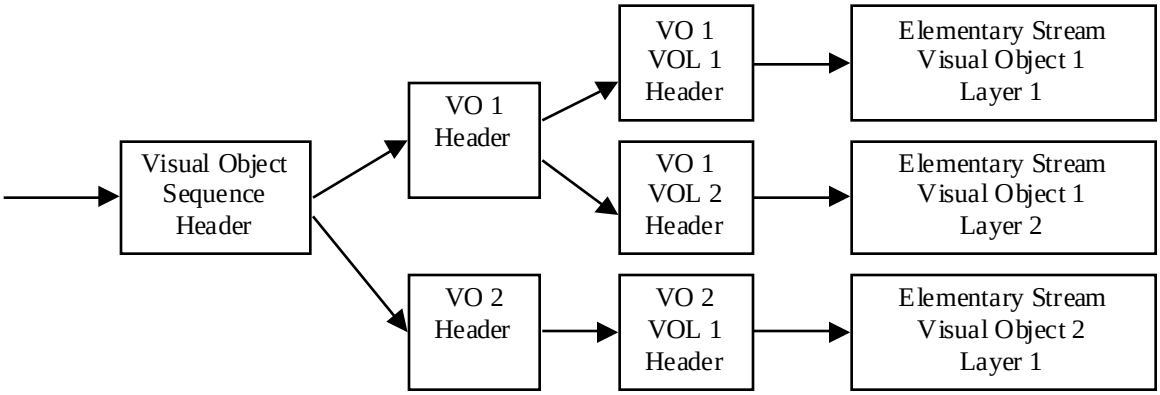


Figure 6-11 -- Example Visual Information – Logical Structure

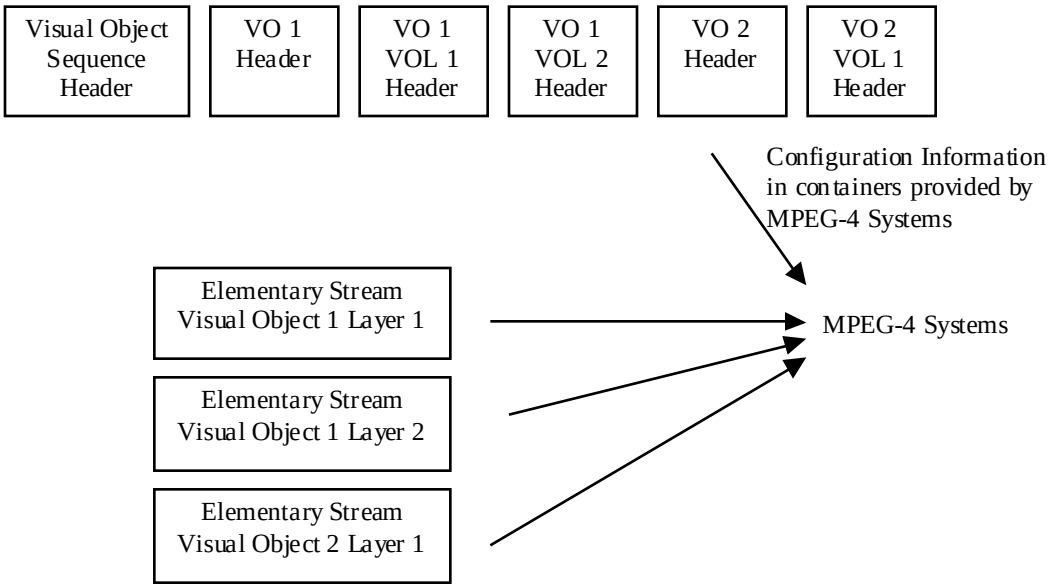


Figure 6-12 -- Example Visual Bitstream – Separate Configuration Information / Elementary Stream.

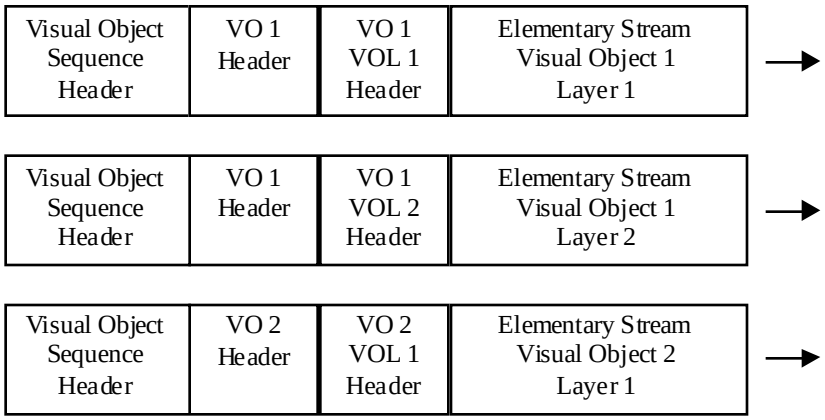


Figure 6-13 -- Example Visual Bitstream – Combined Configuration Information / Elementary Stream

The following functions are entry points for elementary streams, and entry into these functions defines the breakpoint between configuration information and elementary streams:

1. Group_of_VideoObjectPlane(),

2. VideoObjectPlane(),
3. video_plane_with_short_header(),
4. MeshObject(),
5. FaceObject().

FBA Object

fba_object() {	**No. of bits**	**Mnemonic**
fba_object_start_code	32	bslbf
do {		
fba_object_plane()		
} while (!((nextbits_bytealigned() == '000 0000 0000 0000 0000 0000') && (nextbits_bytealigned() != **fba**_object_plane_start_code)))		
}		

FBA Object Plane

fba_object_plane() {	**No. of bits**	**Mnemonic**
fba_object_plane_header()		
fba_object_plane_data()		
}		

fba_object_plane_header() {	**No. of bits**	**Mnemonic**
if (nextbits_bytealigned()=='000 0000 0000 0000 0000 0000'){		
next_start_code()		
fba_object_plane_start_code	32	bslbf
}		
is_intra	1	bslbf
fba_object_mask	2	bslbf
temporal_header()		
}		

fba_object_plane_data() {	**No. of bits**	**Mnemonic**
if(fba_object_mask &'01') {		
if(is_intra) {		
fap_quant	5	uimsbf
for (group_number = 1; group_number <= 10; group_number++) {		
marker_bit	1	uimsbf
fap_mask_type	2	bslbf

if(fap_mask_type == ‘01’|| fap_mask_type == ‘10’)		
fap_group_mask[group_number]	2-16	vlcbf
}		
fba_suggested_gender	1	bslbf
fba_object_coding_type	1	bslbf
if(fba_object_coding_type == 0) {		
is_i_new_max	1	bslbf
is_i_new_min	1	bslbf
is_p_new_max	1	bslbf
is_p_new_min	1	bslbf
decode_new_minmax()		
decode_ifap()		
}		
if(fba_object_coding_type == 1)		
decode_i_segment()		
}		
else {		
if(fba_object_coding_type == 0)		
decode_pfap()		
if(fba_object_coding_type == 1)		
decode_p_segment()		
}		
}		
}		

if(fba_object_mask &'10') {		
if(is_intra) {		
bap_pred_quant_index	5	uimsbf
for (group_number = 1; group_number <= BAP_NUM_GROUPS; group_number++) {		
marker_bit	1	uimsbf
bap_mask_type	2	bslbf
if(bap_mask_type == '01')		
bap_group_mask[group_number]	3-22	vlcbf
else if (bap_mask_type == '00') {		
for(i=0; i<BAPS_IN_GROUP[group_number];i++) {		
bap_group_mask[group_mask][i] = 0		
}		
}		
else if (bap_mask_type == '11') {		
for(i=0; i<BAPS_IN_GROUP[group_number];i++) {		
bap_group_mask[group_mask][i] = 1		
}		
}		
}		
fba_suggested_gender	1	bslbf
fba_object_coding_type	1	bslbf
if (fba_object_coding_type == 0) {		
bap_is_i_new_max	1	bslbf
bap_is_i_new_min	1	bslbf
bap_is_p_new_max	1	bslbf
bap_is_p_new_min	1	bslbf
decode_bap_new_minmax()		
decode_bap_ibap()		
}		
if(fba_object_coding_type == 1)		
decode_bap_i_segment()		
}		
else {		
if (fba_object_coding_type == 0)		
decode_bap_pbap()		
if(fba_object_coding_type == 1)		
decode_bap_p_segment()		
}		
}		

temporal_header() {	No. of bits	Mnemonic
if (is_intra) {		
if (fba_object_mask & '01') {		
is_frame_rate	1	bslbf
if(is_frame_rate)		
decode_frame_rate()		
is_time_code	1	bslbf
if (is_time_code)		
time_code	18	bslbf
}		
}		
skip_frames	1	bslbf
if(skip_frames)		
decode_skip_frames()		
}		

Decode frame rate and skip frames

decode_frame_rate(){	No. of bits	Mnemonic
frame_rate	8	uimsbf
seconds	4	uimsbf
frequency_offset	1	uimsbf
}		

decode_skip_frames(){	No. of bits	Mnemonic
do{		
number_of_frames_to_skip	4	uimsbf
} while (number_of_frames_to_skip = "1111")		
}		

Decode new minmax

decode_new_minmax() {	No. of bits	Mnemonic
if (is_i_new_max) {		
for (group_number = 2, j=0, group_number <= 10, group_number++)		
for (i=0; i < NFAP[group_number]; i++, j++) {		
if (!(i & 0x3))		
marker_bit	1	uimsbf
if (fap_group_mask[group_number] & (1 <<i))		
i_new_max[j]	5	uimsbf

}		
if (is_i_new_min) {		
for (group_number = 2, j=0, group_number <= 10, group_number++)		
for (i=0; i < NFAP[group_number]; i++, j++) {		
if (!(i & 0x3))		
marker_bit	1	uimsbf
if (fap_group_mask[group_number] & (1 <<i))		
i_new_min[j]	5	uimsbf
}		
if (is_p_new_max) {		
for (group_number = 2, j=0, group_number <= 10, group_number++)		
for (i=0; i < NFAP[group_number]; i++, j++) {		
if (!(i & 0x3))		
marker_bit	1	uimsbf
if (fap_group_mask[group_number] & (1 <<i))		
p_new_max[j]	5	uimsbf
}		
if (is_p_new_min) {		
for (group_number = 2, j=0, group_number <= 10, group_number++)		
for (i=0; i < NFAP[group_number]; i++, j++) {		
if (!(i & 0x3))		
marker_bit	1	uimsbf
if (fap_group_mask[group_number] & (1 <<i))		
p_new_min[j]	5	uimsbf
}		
}		
}		

Decode ifap

decode_ifap(){	**No. of bits**	**Mnemonic**
for (group_number = 1, j=0; group_number <= 10; group_number++) {		
if (group_number == 1) {		
if(fap_group_mask[1] & 0x1)		
decode_viseme()		
if(fap_group_mask[1] & 0x2)		
decode_expression()		
} else {		
for (i= 0; i<NFAP[group_number]; i++, j++) {		
if(fap_group_mask[group_number] & (1 << i)) {		
aa_decode(ifap_Q[j],ifap_cum_freq[j])		

}		
}		
}		
}		
}		

Decode pfap

decode_pfap(){	**No. of bits**	**Mnemonic**
for (group_number = 1, j=0; group_number <= 10; group_number++) {		
if (group_number == 1) {		
if(fap_group_mask[1] & 0x1)		
decode_viseme()		
if(fap_group_mask[1] & 0x2)		
decode_expression()		
} else {		
for (i= 0; i<NFAP[group_number]; i++, j++) {		
if(fap_group_mask[group_number] & (1 << i)) {		
aa_decode(pfap_diff[j], pfap_cum_freq[j])		
}		
}		
}		
}		
}		

Decode viseme and expression

decode_viseme() {	**No. of bits**	**Mnemonic**
aa_decode(viseme_select1Q, viseme_select1_cum_freq)		vlclbf
aa_decode(viseme_select2Q, viseme_select2_cum_freq)		vlclbf
aa_decode(viseme_blendQ, viseme_blend_cum_freq)		vlclbf
viseme_def	1	bslbf
}		

decode_expression() {	**No. of bits**	**Mnemonic**
aa_decode(expression_select1Q, expression_select1_cum_freq)		vlclbf
aa_decode(expression_intensity1Q, expression_intensity1_cum_freq)		vlclbf
aa_decode(expression_select2Q, expression_select2_cum_freq)		vlclbf
aa_decode(expression_intensity2Q, expression_intensity2_cum_freq)		vlclbf

aa_decode(expression_blendQ, expression_blend_cum_freq)		vlclbf
init_face	1	bslbf
expression_def	1	bslbf
}		

FBA Object Plane Group

fba_object_plane_group() {	No. of bits	Mnemonic
fba_object_plane_start_code	32	bslbf
is_intra	1	bslbf
if (is_intra) {		
fba_object_mask	2	bslbf
if (fba_object_mask & '01') {		
is_frame_rate	1	bslbf
if (is_frame_rate)		
decode_frame_rate()		
is_time_code	1	bslbf
if (is_time_code)		
time_code	18	
}		
if (fba_object_mask & '10') {		
is_bap_frame_rate	1	bslbf
if (is_bap_frame_rate)		
decode_bap_frame_rate()		
is_bap_time_code	1	bslbf
if (is_bap_time_code)		
bap_time_code	18	bslbf
}		
if (skip_frames)		
decode_skip_frames()		
if (fba_object_mask & '01') {		
fap_quant_index	5	uimsbf
for (group_number=1 to 10) {		
marker_bit	1	uimsbf
fap_mask_type	2	bslbf
if (fap_mask_type == '01' || fap_mask_type == '10')		
fap_group_mask[group_number]	2-16	vlcbf
}		
decode_i_segment()		
}		
if (fba_object_mask & '10') {		
bap_quant_index	5	uimsbf
for (group_number = 1 to BAP_NUM_GROUPS) {		
marker_bit	1	uimsbf
bap_mask_type	2	bslbf

if(bap_mask_type == '01')		
bap_group_mask[group_number]	3-22	vlcbf
else if (bap_mask_type == '00') {		
for(i=0; i<BAPS_IN_GROUP[group_number];i++) {		
bap_group_mask[group_mask][i] = 0		
}		
}		
else if (bap_mask_type == '11') {		
for (i=0; i<BAPS_IN_GROUP[group_number];i++) {		
bap_group_mask[group_mask][i] = 1		
}		
}		
}		
decode_bap_i_segment()		
} else {		
fba_object_group_prediction()		
}		
next_start_code()		
}		

Face Object Group Prediction

fba_object_group_prediction() {	**No. of bits**	**Mnemonic**
skip_frames	1	bslbf
if(skip_frames)		
decode_skip_frames()		
if(fba_object_mask & '01') {		
decode_p_segment()		
}		
if(fba_object_mask & '10') {		
decode_bap_p_segment()		
}		
}		

Decode i_segment

decode_i_segment(){	**No. of bits**	**Mnemonic**
for (group_number= 1, j=0; group_number<= 10; group_number++) {		
if (group_number == 1) {		
if(fap_group_mask[1] & 0x1)		
decode_i_viseme_segment()		

if(fap_group_mask[1] & 0x2)		
decode_i_expression_segment()		
} else {		
for(i=0; i<NFAP[group_number]; i++, j++) {		
if(fap_group_mask[group_number] & (1 << i)) {		
decode_i_dc(dc_Q[j])		
decode_ac(ac_Q[j])		
}		
}		
}		
}		
}		

Decode p_segment

decode_p_segment(){	**No. of bits**	**Mnemonic**
for (group_number = 1, j=0; group_number <= 10; group_number++) {		
if (group_number == 1) {		
if(fap_group_mask[1] & 0x1)		
decode_p_viseme_segment()		
if(fap_group_mask[1] & 0x2)		
decode_p_expression_segment()		
} else {		
for (i=0; i<NFAP[group_number]; i++, j++) {		
If(fap_group_mask[group_number] & (1 << i)) {		
decode_p_dc(dc_Q[j])		
decode_ac(ac_Q[j])		
}		
}		
}		
}		
}		

Decode viseme and expression

decode_i_viseme_segment(){	**No. of bits**	**Mnemonic**
viseme_segment_select1q[0]	4	uimsbf
viseme_segment_select2q[0]	4	uimsbf
viseme_segment_blendq[0]	6	uimsbf
viseme_segment_def[0]	1	bslbf
for (k=1; k<16, k++) {		

viseme_segment_select1q_diff[k]		vlclbf
viseme_segment_select2q_diff[k]		vlclbf
viseme_segment_blendq_diff[k]		vlclbf
viseme_segment_def[k]	1	bslbf
}		
}		

decode_p _viseme_segment(){	**No. of bits**	**Mnemonic**
for (k=0; k<16, k++) {		
viseme_segment_select1q_diff[k]		vlclbf
viseme_segment_select2q_diff[k]		vlclbf
viseme_segment_blendq_diff[k]		vlclbf
viseme_segment_def[k]	1	bslbf
}		
}		

decode_i_expression_segment(){	**No. of bits**	**Mnemonic**
expression_segment_select1q[0]	4	uimsbf
expression_segment_select2q[0]	4	uimsbf
expression_segment_intensity1q[0]	6	uimsbf
expression_segment_intensity2q[0]	6	uimsbf
expression_segment_init_face[0]	1	bslbf
expression_segment_def[0]	1	bslbf
for (k=1; k<16, k++) {		
expression_segment_select1q_diff[k]		vlclbf
expression_segment_select2q_diff[k]		vlclbf
expression_segment_intensity1q_diff[k]		vlclbf
expression_segment_intensity2q_diff[k]		vlclbf
expression_segment_init_face[k]	1	bslbf
expression_segment_def[k]	1	bslbf
}		
}		

decode_p _expression_segment(){	**No. of bits**	**Mnemonic**
for (k=0; k<16, k++) {		
expression_segment_select1q_diff[k]		vlclbf
expression_segment_select2q_diff[k]		vlclbf
expression_segment_intensity1q_diff[k]		vlclbf
expression_segment_intensity2q_diff[k]		vlclbf
expression_segment_init_face[k]	1	bslbf

expression_segment_def[k]	1	bslbf
}		
}		

decode_i_dc(dc_q) {	**No. of bits**	**Mnemonic**
dc_q	16	simsbf
if(dc_q == -256*128)		
dc_q	31	simsbf
}		

decode_p_dc(dc_q_diff) {	**No. of bits**	**Mnemonic**
dc_q_diff		vlclbf
dc_q_diff = dc_q_diff- 256		
if(dc_q_diff == -256)		
dc_q_diff	16	simsbf
if(dc_Q == 0-256*128)		
dc_q_diff	32	simsbf
}		

decode_ac(ac_Q[i]) {	**No. of bits**	**Mnemonic**
this = 0		
next = 0		
while(next < 15) {		
count_of_runs		vlclbf
if (count_of_runs == 15)		
next = 16		
else {		
next = this+1+count_of_runs		
for (n=this+1; n<next; n++)		
ac_q[i][n] = 0		
ac_q[i][next]		vlclbf
if(ac_q[i][next] == 256)		
decode_i_dc(ac_q[i][next])		
else		
ac_q[i][next] = ac_q[i][next]-256		
this = next		
}		
}		
}		

Decode bap min max

decode_bap_new_minmax() {	**No. of bits**	**Mnemonic**
if (bap_is_i_new_max) {		
for (group_number=1;group_number<= BAP_NUM_GROUPS; group_number++)		
for (i=0; i < NBAP_GROUP[group_number]; i++) {		
j=BAPS_IN_GROUP[group_number][i]		
if (!(i & 0x3))		
marker_bit	1	uimsbf
if (bap_group_mask[group_number] & (1 <<i))		
bap_i_new_max[j]	5	uimsbf
}		
if (bap_is_i_new_min) {		
for (group_number = 1; group_number <= BAP_NUM_GROUPS; group_number++)		
for (i=0; i < NBAP_GROUP[group_number]; i++) {		
j=BAPS_IN_GROUP[group_number][i]		
if (!(i & 0x3))		
marker_bit	1	uimsbf
if (bap_group_mask[group_number] & (1 <<i))		
bap_i_new_min[j]	5	uimsbf
}		
if (bap_is_p_new_max) {		
for (group_number = 1; group_number <= BAP_NUM_GROUPS; group_number++)		
for (i=0; i < NBAP_GROUP[group_number]; i++) {		
j=BAPS_IN_GROUP[group_number][i]		
if (!(i & 0x3))		
marker_bit	1	uimsbf
if (bap_group_mask[group_number] & (1 <<i))		
bap_p_new_max[j]	5	uimsbf
}		
if (bap_is_p_new_min) {		
for (group_number = 1; group_number <= BAP_NUM_GROUPS; group_number++)		
for (i=0; i < NBAP_GROUP[group_number]; i++) {		
j=BAPS_IN_GROUP[group_number][i]		
if (!(i & 0x3))		
marker_bit	1	uimsbf

if (bap_group_mask[group_number] & (1 <<i))		
bap_p_new_min[j]	5	uimsbf
}		
}		
}		

Decode ibap

decode_ibap(){	**No. of bits**	**Mnemonic**
for (group_number = 1; group_number <= BAP_NUM_GROUPS; group_number++) {		
for (i= 0; i<NBAP_GROUP[group_number]; i++) {		
j=BAPS_IN_GROUP[group_number][i]		
if(bap_group_mask[group_number] & (1 << i)) {		
aa_decode(ibap_Q[j],ibap_cum_freq[j])		
}		
}		
}		
}		

Decode pbap

decode_pbap() {	**No. of bits**	**Mnemonic**
for (group_number = 1; group_number <= BAP_NUM_GROUPS; group_number++) {		
for (i= 0; i<NBAP_GROUP[group_number]; i++) {		
j=BAPS_IN_GROUP[group_number][i]		
if(bap_group_mask[group_number] & (1 << i)) {		
aa_decode(pbap_diff[j], pbap_cum_freq[j])		
}		
}		
}		
}		

Decode bap i segment

decode_bap_i_segment(){	No. of bits	Mnemonic
for (group_number= 1; group_number<= BAP_NUM_GROUPS; group_number++) {		
for(i=0; i<NBAP_GROUP[group_number]; i++) {		
if(bap_group_mask[group_number] & (1 << i)) {		
j=BAPS_IN_GROUP[group_number][i]		
decode_i_dc(dc_Q[j])		
decode_ac(ac_Q[j])		
}		
}		
}		
}		

Decode bap p segment

decode_bap_p_segment(){	No. of bits	Mnemonic
for (group_number = 1; group_number <= BAP_NUM_GROUPS; group_number++) {		
for (i=0; i<NBAP_GROUP[group_number]; i++) {		
If(bap_group_mask[group_number] & (1 << i)) {		
j = BAPS_IN_GROUP[group_number][i]		
decode_p_dc(dc_Q[j])		
decode_ac(ac_Q[j])		
}		
}		
}		
}		

FBA object

fba_object_start_code: The fba_object_start_code is the bit string '000001BA' in hexadecimal. It initiates a FBA object.

fba_object_coding_type: This is a 1-bit integer indicating which coding method is used. Its meaning is described in Table 6-39.

Table 6-39 -- fba_object_coding_type

type value	Meaning
0	predictive coding
1	DCT (face_object_plane_group)

fba_suggested_gender: This is a 1-bit integer indicating the suggested gender for the face model. It does not bind the decoder to display a facial model of suggested gender, but indicates that the content would be more suitable for display

with the facial model of indicated gender, if the decoder can provide one. If fba_suggested_gender is 1, the suggested gender is male, otherwise it is female.

FBA object plane

face_paramset_mask: This is a 2-bit integer defined in Table 6-40. It indicates whether FBA and BAP data are present in the FBA frame.

Table 6-40 – FBA object mask

mask value	Meaning
00	unused
01	FAP present
10	BAP present
11	both FAP and BAP present

fba_object_plane_start_code: The **fba**_frame_start_code is the bit string '000001BB' in hexadecimal. It initiates a FBA object plane.

is_frame_rate: This is a 1-bit flag which when set to '1' indicates that frame rate information follows this bit field. When set to '0' no frame rate information follows this bit field.

is_time_code: This is a 1-bit flag which when set to '1' indicates that time code information follows this bit field. When set to '0' no time code information follows this bit field.

time_code: This is a 18-bit integer containing the following: time_code_hours, time_code_minutes, marker_bit and time_code_seconds as shown in Table 6-41. The parameters correspond to those defined in the IEC standard publication 461 for "time and control codes for video tape recorders". The time code specifies the modulo part (i.e. the full second units) of the time base for the current object plane.

Table 6-41 -- Meaning of time_code

time_code	range of value	No. of bits	Mnemonic
time_code_hours	0 - 23	5	uimsbf
time_code_minutes	0 - 59	6	uimsbf
marker_bit	1	1	bslbf
time_code_seconds	0 - 59	6	uimsbf

is_bap_frame_rate: This is a 1-bit flag which when set to '1' indicates that frame rate information follows this bit field. When set to '0' no frame rate information follows this bit field.

is_bap_time_code: This is a 1-bit flag which when set to '1' indicates that time code information follows this bit field. When set to '0' no time code information follows this bit field.

bap_time_code: This is a 18-bit integer containing the following: time_code_hours, time_code_minutes, marker_bit and time_code_seconds as shown in . The parameters correspond to those defined in the IEC standard publication 461 for "time and control codes for video tape recorders". The time code refers to the first plane (in display order) after the GOV header. Table 6-31 shows the meaning of time_code.

skip_frames: This is a 1-bit flag which when set to '1' indicates that information follows this bit field that indicates the number of skipped frames. When set to '0' no such information follows this bit field.

fap_mask_type: This is a 2-bit integer. It indicates if the group mask will be present for the specified fap group, or if the complete faps will be present; its meaning is described in Table 6-42. In the case the type is '10' the '0' bit in the group mask indicates interpolate fap.

Table 6-42 -- fap mask type

mask type	Meaning
00	no mask nor fap
01	group mask
10	group mask'

11	fap

fap_group_mask[group_number]: This is a variable length bit entity that indicates, for a particular group_number which fap is represented in the bitstream. The value is interpreted as a mask of 1-bit fields. A 1-bit field in the mask that is set to '1' indicates that the corresponding fap is present in the bitstream. When that 1-bit field is set to '0' it indicates that the fap is not present in the bitstream. The number of bits used for the fap_group_mask depends on the group_number, and is given in Table 6-43.

Table 6-43 -- fap group mask bits

group_number	No. of bits
1	2
2	16
3	12
4	8
5	4
6	5
7	3
8	10
9	4
10	4

NFAP[group_number] **:** This indicates the number of FAPs in each FAP group. Its values are specified in the following table:

Table 6-44 -- NFAP definition

group_number	NFAP[group_number]
1	2
2	16
3	12
4	8
5	4
6	5
7	3
8	10
9	4
10	4

fap_quant: This is a 5-bit unsigned integer which is the quantization scale factor used to compute the FAPi table step size.

is_i_new_max: This is a 1-bit flag which when set to '1' indicates that a new set of maximum range values for I frame follows these 4, 1-bit fields.

is_i_new_min: This is a 1-bit flag which when set to '1' indicates that a new set of minimum range values for I frame follows these 4, 1-bit fields.

is_p_new_max: This is a 1-bit flag which when set to '1' indicates that a new set of maximum range values for P frame follows these 4, 1-bit fields.

is_p_new_min: This is a 1-bit flag which when set to '1' indicates that a new set of minimum range values for P frame follows these 4, 1-bit fields.

bap_group_mask: This mask specifies which BAP groups are to be transmitted. The value is interpreted as a mask of 1-bit fields. A 1-bit field in the mask that is set to '1' indicates that the corresponding group is present in the bitstream. When that 1-bit field is set to '0' it indicates that the bap group is not present in the bitstream. The number of bits used for the bap_group_mask depends on the group_number, and is given below.

bap_pred_quant_index: This is a 5-bit unsigned integer used as the index to a bap_pred_scale table for computing the quantisation step size of BAP values for predictive coding. The value of bap_pred_scale is specified in the following list:

bap_pred_quant_index[0 - 31] =
{ 0, 1, 2, 3, 5, 7, 9, 11, 14, 17, 20, 23, 27, 31, 35, 39, 43, 47, 52, 57, 62, 67, 72, 77, 82, 88, 94, 100, 106, 113, 120, 127}

bap_is_i_new_max: This is a 1-bit flag which when set to '1' indicates that a new set of maximum range values for I frame follows these 4, 1-bit fields.

bap_is_i_new_min: This is a 1-bit flag which when set to '1' indicates that a new set of minimum range values for I frame follows these 4, 1-bit fields.

bap_is_p_new_max: This is a 1-bit flag which when set to '1' indicates that a new set of maximum range values for P frame follows these 4, 1-bit fields.

bap_is_p_new_min: This is a 1-bit flag which when set to '1' indicates that a new set of minimum range values for P frame follows these 4, 1-bit fields.

bap_mask_type: This 2-bit value determines whether BAPs are transmitted individually or in groups.

bap_mask_type	**Meaning**
00	No BAPs
01	BAPs transmitted in groups
10	reserved
11	BAPs transmitted individually

bap_group_mask: this is a variable-length mask indicating which BAPs in a group are present in the fba_object_plane.

group number	group name	No. of. bits
1	Pelvis	3
2	Left leg1	4
3	Right leg1	4
4	Left leg2	6
5	Right leg2	6
6	Left arm1	5
7	Right arm1	5
8	Left arm2	7
9	Right arm2	7
10	Spine1	12
11	Spine2	15
12	Spine3	18
13	Spine4	18
14	Spine5	12
15	Left hand1	16
16	Right hand1	16
17	Left hand2	13
18	Right hand2	13
19	Global positioning	6
20	Extension1	22
21	Extension2	22
22	Extension3	22
23	Extension4	22
24	Extension5	22

Face Object Prediction

skip_frames: This is a 1-bit flag which when set to '1' indicates that information follows this bit field that indicates the number of skipped frames. When set to '0' no such information follows this bit field.

Decode frame rate and frame skip

frame_rate: This is an 8 bit unsigned integer indicating the reference frame rate of the sequence.
seconds: This is a 4 bit unsigned integer indicating the fractional reference frame rate. The frame rate is computed as follows frame rate = (frame_rate + seconds/16).
frequency_offset: This is a 1-bit flag which when set to '1' indicates that the frame rate uses the NTSC frequency offset of 1000/1001. This bit would typically be set when frame_rate = 24, 30 or 60, in which case the resulting frame rate would be 23.97, 29.94 or 59.97 respectively. When set to '0' no frequency offset is present. I.e. if (frequency_offset ==1) frame rate = (1000/1001) * (frame_rate + seconds/16).
number_of_frames_to_skip: This is a 4-bit unsigned integer indicating the number of frames skipped. If the number_of_frames_to skip is equal to 15 (pattern "1111") then another 4-bit word follows allowing to skip up to 29 frames(pattern "11111110"). If the 8-bits pattern equals "11111111", then another 4-bits word will follow and so on, and the number of frames skipped is incremented by 30. Each 4-bit pattern of '1111' increments the total number of frames to skip with 15.
bap_frame_rate: This is an 8 bit unsigned integer indicating the reference frame rate of the face sequence.

bap_seconds: This is a 4 bit unsigned integer indicating the fractional reference frame rate for the face. The frame rate is computed as follows frame rate = (frame_rate + seconds/16).
bap_frequency_offset: This is a 1-bit flag which when set to '1' indicates that the frame rate uses the NTSC frequency offset of 1000/1001. This bit would typically be set when frame_rate = 24, 30 or 60, in which case the resulting frame rate would be 23.97, 29.94 or 59.97 respectively. When set to '0' no frequency offset is present. I.e. if (frequency_offset ==1) frame rate = (1000/1001) * (frame_rate + seconds/16).

Decode new minmax

i_new_max[j]: This is a 5-bit unsigned integer used to scale the maximum value of the arithmetic decoder used in the I frame.
i_new_min[j]: This is a 5-bit unsigned integer used to scale the minimum value of the arithmetic decoder used in the I frame.
p_new_max[j]: This is a 5-bit unsigned integer used to scale the maximum value of the arithmetic decoder used in the P frame.
p_new_min[j]: This is a 5-bit unsigned integer used to scale the minimum value of the arithmetic decoder used in the P frame.

Decode viseme and expression

viseme_def: This is a 1-bit flag which when set to '1' indicates that the mouth FAPs sent with the viseme FAP may be stored in the decoder to help with FAP interpolation in the future.
expression_def: This is a 1-bit flag which when set to '1' indicates that the FAPs sent with the expression FAP may be stored in the decoder to help with FAP interpolation in the future.

FBA object plane group

fba_object_plane_start_code: Defined in subclause 6.3.10.1.
is_intra: This is a 1-bit flag which when set to '1' indicates that the FBA object is coded in intra mode. When set to '0' it indicates that the FBA object is coded in predictive mode.
fba_object_mask: Defined in subclause 6.3.10.1.
is_frame_rate: Defined in subclause 6.3.10.1.
is_time_code: Defined in subclause 6.3.10.1.
time_code: Defined in subclause 6.3.10.1.
skip_frames: Defined in subclause 6.3.10.1.
Fap_quant_index: This is a 5-bit unsigned integer used as the index to a fap_scale table for computing the quantization step size of DCT coefficients. The value of fap_scale is specified in the following list:

fap_scale[0 - 31] = { 1, 1, 2, 3, 5, 7, 8, 10, 12, 15, 18, 21, 25, 30, 35, 42,
50, 60, 72, 87, 105, 128, 156, 191, 234, 288, 355, 439, 543, 674, 836, 1039}

fap_mask_type: Defined in subclause 6.3.10.1.
fap_group_mask[group_number] : Defined in subclause 6.3.10.1.
is_bap_frame_rate: Defined in subclause 6.3.11.1.
is_bap_time_code: Defined in subclause 6.3.11.1.
bap_time_code: Defined in subclause 6.3.11.1.
bap_quant_index: This is a 5-bit unsigned integer used as the index to a bap_scale table for computing the quantisation step size of DCT coefficients. The value of bap_scale is specified in the following list:

bap_scale[0 - 31] = { 1, 1, 2, 3, 5, 7, 8, 10, 12, 15, 18, 21, 25, 30, 35, 42,
50, 60, 72, 87, 105, 128, 156, 191, 234, 288, 355, 439, 543, 674, 836, 1039}

bap_mask_type: Defined in subclause 6.3.11.1
bap_group_mask[group_number]**:** Defined in subclause 6.3.11.1

Face Object Group Prediction

skip_frames: See the definition in subclause 6.3.10.1.

Decode frame rate and frame skip

frame_rate: See the definition in subclause 6.3.10.3.
frequency_offset: See the definition in subclause 6.3.10.3.
number_of_frames_to_skip: See the definition in subclause 6.3.10.3.

Decode viseme_segment and expression_segment

viseme_segment_select1q[k]: This is the quantized value of viseme_select1 at frame k of a viseme FAP segment.

viseme_segment_select2q[k]: This is the quantized value of viseme_select2 at frame k of a viseme FAP segment.
viseme_segment_blendq[k]: This is the quantized value of viseme_blend at frame k of a viseme FAP segment.
viseme_segment_def[k]: This is a 1-bit flag which when set to '1' indicates that the mouth FAPs sent with the viseme FAP at frame k of a viseme FAP segment may be stored in the decoder to help with FAP interpolation in the future.
viseme_segment_select1q_diff[k]: This is the prediction error of viseme_select1 at frame k of a viseme FAP segment.
viseme_segment_select2q_diff[k]: This is the prediction error of viseme_select2 at frame k of a viseme FAP segment.
viseme_segment_blendq_diff[k]: This is the prediction error of viseme_blend at frame k of a viseme FAP segment.
expression_segment_select1q[k]: This is the quantized value of expression_select1 at frame k of an expression FAP segment.
expression_segment_select2q[k]: This is the quantized value of expression_select2 at frame k of an expression FAP segment.
expression_segment_intensity1q[k]: This is the quantized value of expression_intensity1 at frame k of an expression FAP segment
expression_segment_intensity2q[k]: This is the quantized value of expression_intensity2 at frame k of an expression FAP segment
expression_segment_select1q_diff[k]: This is the prediction error of expression_select1 at frame k of an expression FAP segment.
expression_segment_select2q_diff[k]: This is the prediction error of expression_select2 at frame k of an expression FAP segment.
expression_segment_intensity1q_diff[k]: This is the prediction error of expression_intensity1 at frame k of an expression FAP segment.
expression_segment_intensity2q_diff[k]: This is the prediction error of expression_intensity2 at frame k of an expression FAP segment.
expression_segment_init_face[k]: This is a 1-bit flag which indicates the value of init_face at frame k of an expression FAP segment.
expression_segment_def[k]: This is a 1-bit flag which when set to '1' indicates that the FAPs sent with the expression FAP at frame k of a viseme FAP segment may be stored in the decoder to help with FAP interpolation in the future.

Decode i_dc, p_dc, and ac

dc_q: This is the quantized DC component of the DCT coefficients. For an intra FAP segment, this component is coded as a signed integer of either 16 bits or 31 bits. The DCT quantization parameters of the 68 FAPs are specified in the following list:

DCTQP[1 - 68] = {1, 1, 7.5, 7.5, 7.5, 7.5, 7.5, 7.5, 7.5, 7.5,
7.5, 7.5, 7.5, 15, 15, 15, 15, 5, 10, 10,
10, 10, 425, 425, 425, 425, 5, 5, 5, 5,
7.5, 7.5, 7.5, 7.5, 7.5, 7.5, 7.5, 7.5, 20, 20,
20, 20, 10, 10, 10, 10, 255, 170, 255, 255,
7.5, 7.5, 7.5, 7.5, 7.5, 7.5, 7.5, 7.5, 7.5, 7.5,
15, 15, 15, 15, 10, 10, 10, 10}

For DC coefficients, the quantization stepsize is obtained as follows:

qstep[i] = fap_scale[fap_quant_inex] * DCTQP[i] ÷ 3.0

dc_q_diff: This is the quantized prediction error of a DC coefficient of an inter FAP segment. Its value is computed by subtracting the decoded DC coefficient of the previous FAP segment from the DC coefficient of the current FAP segment. It is coded by a variable length code if its value is within [-255, +255]. Outside this range, its value is coded by a signed integer of 16 or 32 bits.
count_of_runs: This is the run length of zeros preceding a non-zero AC coefficient.
ac_q[i][next]: This is a quantized AC coefficients of a segment of FAPi. For AC coefficients, the quantization stepsize is three times larger than the DC quantization stepsize and is obtained as follows:

qstep[i] = fap_scale[fap_quant_inex] * DCTQP[i]

Decode bap min max

bap_i_new_max[i] – This is a 5-bit unsigned integer used to scale the maximum value of the arithmetic decoder used in the I frame.
bap_i_new_min[i] – This is a 5-bit unsigned integer used to scale the minimum value of the arithmetic decoder used in the I frame.

bap_p_new_max[i] – This is a 5-bit unsigned integer used to scale the maximum value of the arithmetic decoder used in the P frame.
bap_p_new_min[i] – This is a 5-bit unsigned integer used to scale the minimum value of the arithmetic decoder used in the P frame.

FBA object decoding

Frame based face object decoding

This subclause specifies the additional decoding process required for face object decoding.

The coded data is decoded by an arithmetic decoding process. The arithmetic decoding process is described in detail in annex B. Following the arithmetic decoding, the data is de-quantized by an inverse quantization process. The FAPs are obtained by a predictive decoding scheme as shown in Figure 7-47.
The base quantization step size QP for each FAP is listed in Table C-1. The quantization parameter fap_quant is applied uniformly to all FAPs. The magnitude of the quantization scaling factor ranges from 1 to 8. The value of fap_quant == 0 has a special meaning, it is used to indicate lossless coding mode, so no dequantization is applied. The quantization stepsize is obtained as follows:

```
if (fap_quant)
    qstep = QP * fap_quant
else
    qstep = 1
```

The dequantized FAP'(t) is obtained from the decoded coefficient FAP''(t) as follows:
FAP'(t) = qstep * FAP''(t)

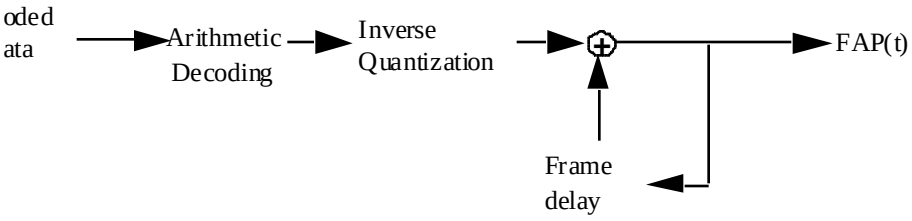


Figure 7-47 -- FAP decoding

Decoding of faps

For a given frame FAPs in the decoder assume one of three of the following states:

1. set by a value transmitted by the encoder
2. retain a value previously sent by the encoder
3. interpolated by the decoder

FAP values which have been initialized in an intra coded FAP set are assumed to retain those values if subsequently masked out unless a special mask mode is used to indicate interpolation by the decoder. FAP values which have never been initialized must be estimated by the decoder. For example, if only FAP group 2 (inner lip) is used and FAP group 8 (outer lip) is never used, the outer lip points must be estimated by the decoder. In a second example the FAP decoder is also expected to enforce symmetry when only the left or right portion of a symmetric FAP set is received (e.g. if the left eye is moved and the right eye is subject to interpolation, it is to be moved in the same way as the left eye).

DCT based face object decoding

The bitstream is decoded into segments of FAPs, where each segment is composed of a temporal sequence of 16 FAP object planes. The block diagram of the decoder is shown in Figure 7-48.

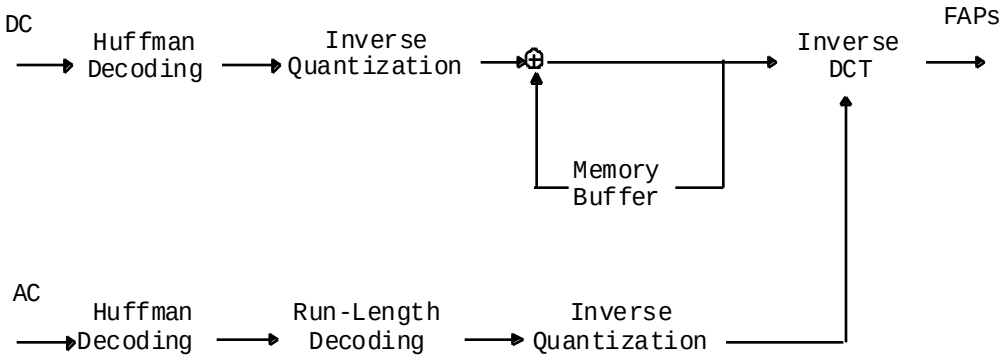


Figure 7-48 -- **Block diagram of the DCT-based decoding process**

The DCT-based decoding process consists of the following three basic steps:

1. Differential decoding the DC coefficient of a segment.
2. Decoding the AC coefficients of the segment
3. Determining the 16 FAP values of the segment using inverse discrete cosine transform (IDCT).

A uniform quantization step size is used for all AC coefficients. The quantization step size for AC coefficients is obtained as follows:

qstep[i] = fap_scale[fap_quant_inex] * DCTQP[i]

where DCTQP[i] is the base quantization step size and its value is defined in subclause 6.3.10.10. The quantization step size of the DC coefficient is one-third of the AC coefficients. Different quantization step sizes are used for different FAPs.

The DCT-based decoding process is applied to all FAP segments except the viseme (FAP #1) and expression (FAP #2) parameters. The latter two parameters are differential decoded without transform. The decoding of viseme and expression segments are described at the end of this subclause.

For FAP #3 to FAP #68, the DC coefficient of an intra coded segment is stored as a 16-bit signed integer if its value is within the 16-bit range. Otherwise, it is stored as a 31-bit signed integer. For an inter coded segment, the DC coefficient of the previous segment is used as a prediction of the current DC coefficient. The prediction error is decoded using a Huffman table of 512 symbols. . An "ESC" symbol, if obtained, indicates that the prediction error is out of the range [-255, 255]. In this case, the next 16 bits extracted from the bitstream are represented as a signed 16-bit integer for the prediction error. If the value of the integer is equal to -256*128, it means that the value of the prediction error is over the 16-bit range. Then the following 32 bits from the bitstream are extracted as a signed 32-bit integer, in twos complement format and the most significant bit first

The AC coefficients, for both inter and intra coded segments, are decoded using Huffman tables. The run-length code indicates the number of leading zeros before each non-zero AC coefficient. The run-length ranges from 0 to 14 and proceeds the code for the AC coefficient. The symbol 15 in the run length table indicates the end of non-zero symbols in a segment. Therefore, the Huffman table of the run-length codes contains 16 symbols. The values of non-zero AC coefficients are decoded in a way similar to the decoding of DC prediction errors but with a different Huffman table.

The bitstreams corresponding to viseme and expression segments are basically differential decoded without IDCT. For an intra coded segment, the quantized values of the first viseme_select1, viseme_select2, viseme_blend, expression_select1, expression_select2, expression_intensity1, and expression_intensity2 within the segment are decoded using fixed length code. These first values are used as the prediction for the second viseme_select1, viseme_select2, … etc of the segment and the prediction error are differential decoded using Huffman tables. For an inter coded segment, the last viseme_select1, for example, of the previous decoded segment is used to predict the first viseme_select1 of the current segment. In general, the decoded values (before inverse quantization) of differential coded viseme and expression parameter fields are obtained

byviseme_segment_select1q[k] = viseme_segment_select1q[k-1] +
viseme_segment_select1q_diff[k] - 14
viseme_segment_select2q[k] = viseme_segment_select2q[k-1] +
viseme_segment_select2q_diff[k] - 14
viseme_segment_blendq[k] = viseme_segment_blendq[k-1] +
viseme_segment_blendq_diff[k] - 63
expression_segment_select1q[k] = expression_segment_select1q[k-1] +
expression_segment_select1q_diff[k] - 6
expression_segment_select2q[k] = expression_segment_select2q[k-1] +
expression_segment_select2q_diff[k] - 6
expression_segment_intensity1q[k] = expression_segment_intensity1q[k-1] +
expression_segment_intensity1q_diff[k] - 63
expression_segment_intensity2q[k] = expression_segment_intensity2q[k-1] +
expression_segment_intensity2q_diff[k] - 63

Decoding of the viseme parameter fap 1

Fourteen visemes have been defined for selection by the Viseme Parameter FAP 1, the definition is given in annex C. The viseme parameter allows two visemes from a standard set to be blended together. The viseme parameter is composed of a set of values as follows.

Table 7-17 -- Viseme parameter range

viseme () {	**Range**
viseme_select1	0-14
viseme_select2	0-14
viseme_blend	0-63
viseme_def	0-1
}	

Viseme_blend is quantized (step size = 1) and defines the blending of viseme1 and viseme2 in the decoder by the following symbolic expression where viseme1 and 2 are graphical interpretations of the given visemes as suggested in the non-normative annex.

final viseme = (viseme 1) * (viseme_blend / 63) + (viseme 2) * (1 - viseme_blend / 63)

The viseme can only have impact on FAPs that are currently allowed to be interpolated.
If the viseme_def bit is set, the current mouth FAPs can be used by the decoder to define the selected viseme in terms of a table of FAPs. This FAP table can be used when the same viseme is invoked again later for FAPs which must be interpolated.

Decoding of the viseme parameter fap 2

The expression parameter allows two expressions from a standard set to be blended together.The expression parameter is composed of a set of values as follows.

Table 7-18 -- Expression parameter range

expression () {	**Range**
expression_select1	0-6
expression_intensity1	0-63
expression_select2	0-6

expression_intensity2	0-63
init_face	0-1
expression_def	0-1
}	

Expression_intensity1 and expression_intensity2 are quantized (step size = 1) and define excitation of expressions 1 and 2 in the decoder by the following equations where expressions 1 and 2 are graphical interpretations of the given expression as suggested by the non-normative reference:

final expression = expression1 * (expression_intensity1 / 63)+ expression2 * (expression_intensity2 / 63)

The decoder displays the expressions according to the above fomula as a superposition of the 2 expressions.
The expression can only have impact on FAPs that are currently allowed to be interpolated. If the init_face bit is set, the neutral face may be modified within the neutral face constraints of mouth closure, eye opening, gaze direction, and head orientation before FAPs 3-68 are applied. If the expression_def bit is set, the current FAPs can be used to define the selected expression in terms of a table of FAPs. This FAP table can then be used when the same expression is invoked again later.

Fap masking

The face is animated by sending a stream of facial animation parameters. FAP masking, as indicated in the bitstream, is used to select FAPs. FAPs are selected by using a two level mask hierarchy. The first level contains two bit code for each group indicating the following options:

1. no FAPs are sent in the group.
2. a mask is sent indicating which FAPs in the group are sent. FAPs not selected by the group mask retain their previous value if any previously set value (not interpolated by decoder if previously set)
3. a mask is sent indicating which FAPs in the group are sent. FAPs not selected by the group mask retain must be interpolated by the decoder.
4. all FAPs in the group are sent.

Frame Based Body Decoding

This clause specifies the additional decoding process required for body model decoding.
The BAPs are quantized and coded by a predictive coding scheme, similar to the FAPs. For each parameter to be coded in the current frame, the decoded value of this parameter in the previous frame is used as the prediction. Then the prediction error, i.e., the difference between the current parameter and its prediction, is computed and coded by arithmetic coding. This predictive coding scheme prevents the coding error from accumulating. The arithmetic decoding process is described in detail in Visual FCD Version 1.
Similar to FAPs, each BAP has a different precision requirement. Therefore different quantisation step sizes are applied to the BAPs. The base quantisation step size for each BAP is defined in the tables below. The bit rate is controlled by adjusting the quantisation step via the use of a quantisation parameter (scaling factor) called BAP_QUANT.
BAP_QUANT is applied uniformly to all BAPs. The magnitude of the quantisation parameter ranges from 0 to 31.
The base quantisation step size BQP for each BAP is listed in Annex C. The quantisation parameter BAP_QUANT is applied uniformly to all BAPs. The magnitude of the quantisation scaling factor ranges from 1 to 31. The value of BAP_QUANT == 0 has a special meaning, it is used to indicate lossless coding mode, so no dequantisation is applied.
The quantisation stepsize is obtained as follows:

```
if (BAP_QUANT)
    qstep = BQP * BAP_QUANT
else
    qstep = 1
```

The dequantized BAP'(t) is obtained from the decoded coefficient BAP''(t) as follows:
BAP'(t) = qstep * BAP''(t)

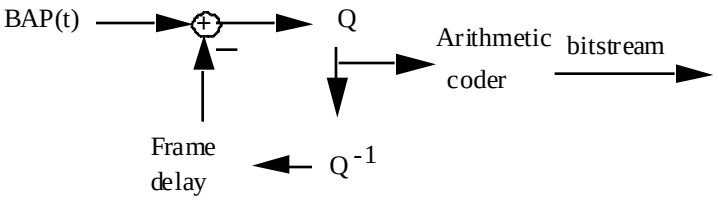


Figure V2 -2 -- BAP predictive coding

Decoding of BAPs

For a given frame, BAPs in the decoder assume one of the three following states:
1. set by a value transmitted by the encoder
2. retain a value previously sent by the encoder
BAP values which have been initialized in an intra coded BAP set are assumed to retain those values if subsequently masked out.

DCT based body object decoding

The bitstream is decoded into segments of BAPs, where each segment is composed of a temporal sequence of 16 BAP object planes. The block diagram of the decoder is shown in Figure V2 -3.

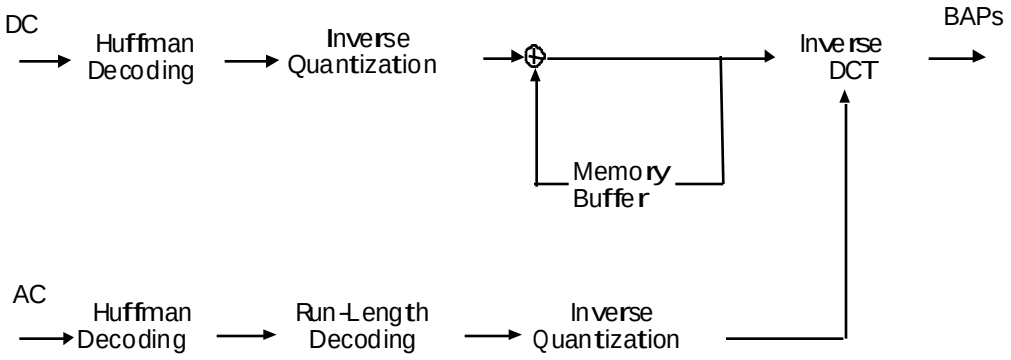


Figure V2 -3 -- Block diagram of the DCT-based BAPs decoding process.

The DCT-based decoding process consists of the following three basic steps:
1. Differential decoding the DC coefficient of a segment.
2. Decoding the AC coefficients of the segment
3. Determining the 16 BAP values of the segment using inverse discrete cosine transform (IDCT).
A uniform quantisation step size is used for all AC coefficients. The quantisation step size for AC coefficients is obtained as follows:

qstep[i] = bap_scale[bap_quant_inex] * BQP[i]

where BQP[i] is the base quantisation step size and its value is defined in Table (TBD). The quantisation step size of the DC coefficient is one-third of the AC coefficients.
The DCT-based decoding process is applied to all BAPs. The DC coefficient of an intra coded segment is stored as a 16-bit signed integer if its value is within the 16-bit range. Otherwise, it is stored as a 31-bit signed integer. For an inter-coded segment, the DC coefficient of the previous segment is used as a prediction of the current DC coefficient. The prediction error is decoded using a Huffman table of 512 symbols. . An "ESC" symbol, if obtained, indicates that the prediction error is out of the range [-255, 255]. In this case, the next 16 bits extracted from the bitstream are represented as a signed 16-bit integer for the prediction error. If the value of the integer is equal to -256*128, it means that the value of the prediction error is over the 16-bit range. Then the following 32 bits from the bitstream are extracted as a signed 32-bit integer, in twos complement format and the most significant bit first
The AC coefficients, for both inter and intra coded segments, are decoded using Huffman tables. The run-length code indicates the number of leading zeros before each non-zero AC coefficient. The run-length ranges from 0 to 14 and proceeds the code for the AC coefficient. The symbol 15 in the run length table indicates the end of non-zero symbols in a segment. Therefore, the Huffman table of the run-length codes contains 16 symbols. The values of non-zero AC coefficients are decoded in a way similar to the decoding of DC prediction errors but with a different Huffman table.

A.1.1 FBA Object Decoding

In FBA decoder, a symbol is decoded by using a specific model based on the syntax and by calling the following procedure which is specified in C.

```
static long low, high, code_value, bit, length, sacindex, cum,
zerorun=0;

int aa_decode(int cumul_freq[ ])
{
  length = high - low + 1;
  cum = (-1 + (code_value - low + 1) * cumul_freq[0]) / length;
  for (sacindex = 1; cumul_freq[sacindex] > cum; sacindex++);
  high = low - 1 + (length * cumul_freq[sacindex-1]) / cumul_freq[0];
  low += (length * cumul_freq[sacindex]) / cumul_freq[0];

  for ( ; ; ) {
    if (high < q2) ;
    else if (low >= q2) {
      code_value -= q2;
      low -= q2;
      high -= q2;
    }
    else if (low >= q1 && high < q3) {
      code_value -= q1;
      low -= q1;
      high -= q1;
    }
    else {
      break;
    }
    low *= 2;
    high = 2*high + 1;
    bit_out_psc_layer();
    code_value = 2*code_value + bit;
    used_bits++;
  }
  return (sacindex-1);
}

void bit_out_psc_layer()
{
  bit = getbits(1);
}
```

Again the model is specified through `cumul_freq[ ]`. The decoded symbol is returned through its index in the model. The decoder is initialized to start decoding an arithmetic coded bitstream by calling the following procedure.

```
void decoder_reset( )
```

```
{
  int i;
  zerorun = 0;        /* clear consecutive zero's counter */
  code_value = 0;
  low = 0;
  high = top;
  for (i = 1;   i <= 16;   i++) {
    bit_out_psc_layer();
    code_value = 2 * code_value + bit;
  }
  used_bits = 0;
}
```

Annex B

(normative)

Face and body object decoding tables and definitions

FAPs names may contain letters with the following meaning: l = left, r = right, t = top, b = bottom, i = inner, o = outer, m = middle. The sum of two corresponding top and bottom eyelid FAPs must equal 1024 when the eyelids are closed. Inner lips are closed when the sum of two corresponding top and bottom lip FAPs equals zero. For example: (lower_t_midlip + raise_b_midlip) = 0 when the lips are closed. All directions are defined with respect to the face and not the image of the face.

Table C-1 -- FAP definitions, group assignments and step sizes

#	FAP name	FAP description	units	Uni-orBidir	Pos motion	Grp	FDP subgrp num	Quant step size	Min/Max I-Frame quantized values	Min/Max P-Frame quantized values
1	viseme	Set of values determining the mixture of two visemes for this frame (e.g. pbm, fv, th)	na	na	na	1	na	1	viseme_blend: +63	viseme_blend: +-63
2	expression	A set of values determining the mixture of two facial expression	na	na	na	1	na	1	expression_intensity1, expression_intensity2: +63	expression_intensity1, expression_intensity2: +-63
3	open_jaw	Vertical jaw displacement (does not affect mouth opening)	MNS	U	down	2	1	4	+1080	+360
4	lower_t_midlip	Vertical top middle inner lip displacement	MNS	B	down	2	2	2	+-600	+-180
5	raise_b_midlip	Vertical bottom middle inner lip displacement	MNS	B	up	2	3	2	+-1860	+-600
6	stretch_l_cornerlip	Horizontal displacement of left inner lip corner	MW	B	left	2	4	2	+-600	+-180
7	stretch_r_cornerlip	Horizontal	MW	B	right	2	5	2	+-600	+-180

		displacement of right inner lip corner								
8	lower_t_lip_lm	Vertical displacement of midpoint between left corner and middle of top inner lip	MNS	B	down	2	6	2	+-600	+-180
9	lower_t_lip_rm	Vertical displacement of midpoint between right corner and middle of top inner lip	MNS	B	down	2	7	2	+-600	+-180
10	raise_b_lip_lm	Vertical displacement of midpoint between left corner and middle of bottom inner lip	MNS	B	up	2	8	2	+-1860	+-600
11	raise_b_lip_rm	Vertical displacement of midpoint between right corner and middle of bottom inner lip	MNS	B	up	2	9	2	+-1860	+-600
12	raise_l_cornerlip	Vertical displacement of left inner lip corner	MNS	B	up	2	4	2	+-600	+-180
13	raise_r_cornerlip	Vertical displacement of right inner lip corner	MNS	B	up	2	5	2	+-600	+-180
14	thrust_jaw	Depth displacement of jaw	MNS	U	forward	2	1	1	+600	+180
15	shift_jaw	Side to side displacement of jaw	MW	B	right	2	1	1	+-1080	+-360
16	push_b_lip	Depth displacement of bottom middle lip	MNS	B	forward	2	3	1	+-1080	+-360
17	push_t_lip	Depth displacement of top middle lip	MNS	B	forward	2	2	1	+-1080	+-360
18	depress_chin	Upward and	MNS	B	up	2	10	1	+-420	+-180

		compressing movement of the chin (like in sadness)								
19	close_t_l_eyelid	Vertical displacement of top left eyelid	IRISD	B	down	3	1	1	+-1080	+-600
20	close_t_r_eyelid	Vertical displacement of top right eyelid	IRISD	B	down	3	2	1	+-1080	+-600
21	close_b_l_eyelid	Vertical displacement of bottom left eyelid	IRISD	B	up	3	3	1	+-600	+-240
22	close_b_r_eyelid	Vertical displacement of bottom right eyelid	IRISD	B	up	3	4	1	+-600	+-240
23	yaw_l_eyeball	Horizontal orientation of left eyeball	AU	B	left	3	na	128	+-1200	+-420
24	yaw_r_eyeball	Horizontal orientation of right eyeball	AU	B	left	3	na	128	+-1200	+-420
25	pitch_l_eyeball	Vertical orientation of left eyeball	AU	B	down	3	na	128	+-900	+-300
26	pitch_r_eyeball	Vertical orientation of right eyeball	AU	B	down	3	na	128	+-900	+-300
27	thrust_l_eyeball	Depth displacement of left eyeball	ES	B	forward	3	na	1	+-600	+-180
28	thrust_r_eyeball	Depth displacement of right eyeball	ES	B	forward	3	na	1	+-600	+-180
29	dilate_l_pupil	Dilation of left pupil	IRISD	B	growing	3	5	1	+-420	+-120
30	dilate_r_pupil	Dilation of right pupil	IRISD	B	growing	3	6	1	+-420	+-120
31	raise_l_i_eyebrow	Vertical displacement of left inner eyebrow	ENS	B	up	4	1	2	+-900	+-360
32	raise_r_i_eyebrow	Vertical displacement of right inner eyebrow	ENS	B	up	4	2	2	+-900	+-360
33	raise_l_m_eyebrow	Vertical displacement of left middle eyebrow	ENS	B	up	4	3	2	+-900	+-360

34	raise_r_m_eyebrow	Vertical displacement of right middle eyebrow	ENS	B	up	4	4	2	+-900	+-360
35	raise_l_o_eyebrow	Vertical displacement of left outer eyebrow	ENS	B	up	4	5	2	+-900	+-360
36	raise_r_o_eyebrow	Vertical displacement of right outer eyebrow	ENS	B	up	4	6	2	+-900	+-360
37	squeeze_l_eyebrow	Horizontal displacement of left eyebrow	ES	B	right	4	1	1	+-900	+-300
38	squeeze_r_eyebrow	Horizontal displacement of right eyebrow	ES	B	left	4	2	1	+-900	+-300
39	puff_l_cheek	Horizontal displacement of left cheeck	ES	B	left	5	1	2	+-900	+-300
40	puff_r_cheek	Horizontal displacement of right cheeck	ES	B	right	5	2	2	+-900	+-300
41	lift_l_cheek	Vertical displacement of left cheek	ENS	U	up	5	3	2	+-600	+-180
42	lift_r_cheek	Vertical displacement of right cheek	ENS	U	up	5	4	2	+-600	+-180
43	shift_tongue_tip	Horizontal displacement of tongue tip	MW	B	right	6	1	1	+-1080	+-420
44	raise_tongue_tip	Vertical displacement of tongue tip	MNS	B	up	6	1	1	+-1080	+-420
45	thrust_tongue_tip	Depth displacement of tongue tip	MW	B	forward	6	1	1	+-1080	+-420
46	raise_tongue	Vertical displacement of tongue	MNS	B	up	6	2	1	+-1080	+-420
47	tongue_roll	Rolling of the tongue into U shape	AU	U	concave upward	6	3, 4	512	+300	+60
48	head_pitch	Head pitch angle from top of spine	AU	B	down	7	na	170	+-1860	+-600

49	head_yaw	Head yaw angle from top of spine	AU	B	left	7	na	170	+-1860	+-600
50	head_roll	Head roll angle from top of spine	AU	B	right	7	na	170	+-1860	+-600
51	lower_t_midlip _o	Vertical top middle outer lip displacement	MNS	B	down	8	1	2	+-600	+-180
52	raise_b_midlip_o	Vertical bottom middle outer lip displacement	MNS	B	up	8	2	2	+-1860	+-600
53	stretch_l_cornerlip_o	Horizontal displacement of left outer lip corner	MW	B	left	8	3	2	+-600	+-180
54	stretch_r_cornerlip_o	Horizontal displacement of right outer lip corner	MW	B	right	8	4	2	+-600	+-180
55	lower_t_lip_lm _o	Vertical displacement of midpoint between left corner and middle of top outer lip	MNS	B	down	8	5	2	+-600	+-180
56	lower_t_lip_rm _o	Vertical displacement of midpoint between right corner and middle of top outer lip	MNS	B	down	8	6	2	+-600	+-180
57	raise_b_lip_lm_o	Vertical displacement of midpoint between left corner and middle of bottom outer lip	MNS	B	up	8	7	2	+-1860	+-600
58	raise_b_lip_rm_o	Vertical displacement of midpoint between right corner and middle of bottom outer lip	MNS	B	up	8	8	2	+-1860	+-600
59	raise_l_cornerlip_o	Vertical displacement of left outer lip corner	MNS	B	up	8	3	2	+-600	+-180
60	raise_r_cornerlip _o	Vertical	MNS	B	up	8	4	2	+-600	+-180

		displacement of right outer lip corner								
61	stretch_l_nose	Horizontal displacement of left side of nose	ENS	B	left	9	1	1	+-540	+-120
62	stretch_r_nose	Horizontal displacement of right side of nose	ENS	B	right	9	2	1	+-540	+-120
63	raise_nose	Vertical displacement of nose tip	ENS	B	up	9	3	1	+-680	+-180
64	bend_nose	Horizontal displacement of nose tip	ENS	B	right	9	3	1	+-900	+-180
65	raise_l_ear	Vertical displacement of left ear	ENS	B	up	10	1	1	+-900	+-240
66	raise_r_ear	Vertical displacement of right ear	ENS	B	up	10	2	1	+-900	+-240
67	pull_l_ear	Horizontal displacement of left ear	ENS	B	left	10	3	1	+-900	+-300
68	pull_r_ear	Horizontal displacement of right ear	ENS	B	right	10	4	1	+-900	+-300

Table C-2 -- FAP grouping

Group	Number of FAPs
1: visemes and expressions	2
2: jaw, chin, inner lowerlip, cornerlips, midlip	16
3: eyeballs, pupils, eyelids	12
4: eyebrow	8
5: cheeks	4
6: tongue	5
7: head rotation	3
8: outer lip positions	10
9: nose	4
10: ears	4

In the following, each facial expression is defined by a textual description and a pictorial example. (reference [10], page 114.) This reference was also used for the characteristics of the described expressions.

Table C-3 -- Values for expression_select

expression_select	expression name	textual description
0	na	na
1	joy	The eyebrows are relaxed. The mouth is open and the mouth corners pulled back toward the ears.
2	sadness	The inner eyebrows are bent upward. The eyes are slightly closed. The mouth is relaxed.
3	anger	The inner eyebrows are pulled downward and together. The eyes are wide open. The lips are pressed against each other or opened to expose the teeth.
4	fear	The eyebrows are raised and pulled together. The inner eyebrows are bent upward. The eyes are tense and alert.
5	disgust	The eyebrows and eyelids are relaxed. The upper lip is raised and curled, often asymmetrically.
6	surprise	The eyebrows are raised. The upper eyelids are wide open, the lower relaxed. The jaw is opened.

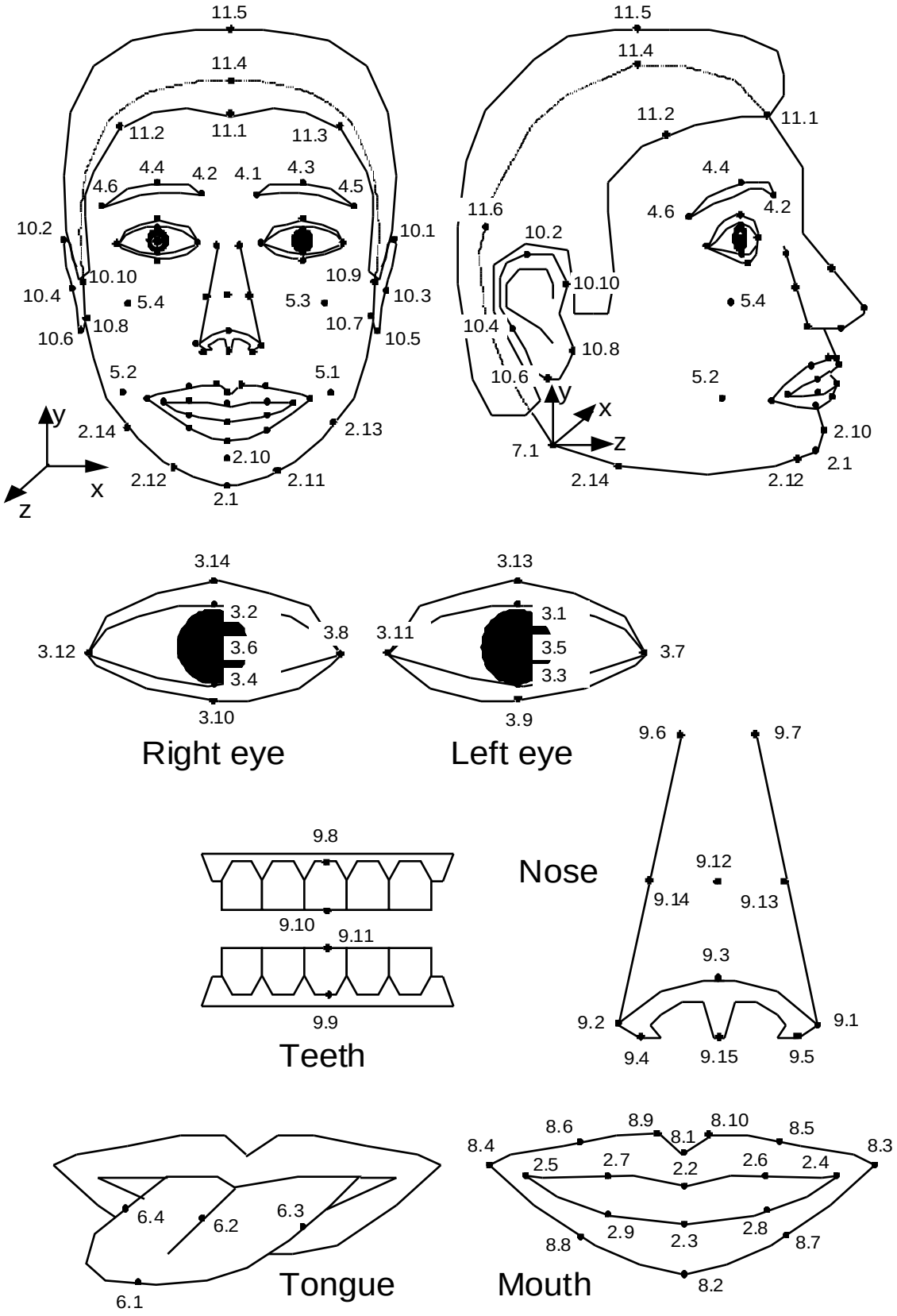


Figure C-1 -- FDP feature point set

In the following, the notation 2.1.x indicates the x coordinate of feature point 2.1.

Feature points		Recommended location constraints		
#	Text description	x	y	z
2.1	Bottom of the chin	7.1.x		
2.2	Middle point of inner upper lip contour	7.1.x		
2.3	Middle point of inner lower lip contour	7.1.x		
2.4	Left corner of inner lip contour			
2.5	Right corner of inner lip contour			
2.6	Midpoint between f.p. 2.2 and 2.4 in the inner upper lip contour	(2.2.x+2.4.x)/2		
2.7	Midpoint between f.p. 2.2 and 2.5 in the inner upper lip contour	(2.2.x+2.5.x)/2		
2.8	Midpoint between f.p. 2.3 and 2.4 in the inner lower lip contour	(2.3.x+2.4.x)/2		
2.9	Midpoint between f.p. 2.3 and 2.5 in the inner lower lip contour	(2.3.x+2.5.x)/2		
2.10	Chin boss	7.1.x		
2.11	Chin left corner	> 8.7.x and < 8.3.x		
2.12	Chin right corner	> 8.4.x and < 8.8.x		
2.13	Left corner of jaw bone			
2.14	Right corner of jaw bone			
3.1	Center of upper inner left eyelid	(3.7.x+3.11.x)/2		
3.2	Center of upper inner right eyelid	(3.8.x+3.12.x)/2		
3.3	Center of lower inner left eyelid	(3.7.x+3.11.x)/2		
3.4	Center of lower inner right eyelid	(3.8.x+3.12.x)/2		
3.5	Center of the pupil of left eye			
3.6	Center of the pupil of right eye			
3.7	Left corner of left eye			
3.8	Left corner of right eye			
3.9	Center of lower outer left eyelid	(3.7.x+3.11.x)/2		
3.10	Center of lower outer right eyelid	(3.7.x+3.11.x)/2		
3.11	Right corner of left eye			
3.12	Right corner of right eye			
3.13	Center of upper outer left eyelid	(3.8.x+3.12.x)/2		
3.14	Center of upper outer right eyelid	(3.8.x+3.12.x)/2		
4.1	Right corner of left eyebrow			
4.2	Left corner of right eyebrow			
4.3	Uppermost point of the left eyebrow	(4.1.x+4.5.x)/2 or x coord of the uppermost point		

		of the contour		
4.4	Uppermost point of the right eyebrow	(4.2.x+4.6.x)/2 or x coord of the uppermost point of the contour		
4.5	Left corner of left eyebrow			
4.6	Right corner of right eyebrow			
5.1	Center of the left cheek		8.3.y	
5.2	Center of the right cheek		8.4.y	
5.3	Left cheek bone	> 3.5.x and < 3.7.x	> 9.15.y and < 9.12.y	
5.4	Right cheek bone	> 3.6.x and < 3.12.x	> 9.15.y and < 9.12.y	
6.1	Tip of the tongue	7.1.x		
6.2	Center of the tongue body	7.1.x		
6.3	Left border of the tongue			6.2.z
6.4	Right border of the tongue			6.2.z
7.1	top of spine (center of head rotation)			
8.1	Middle point of outer upper lip contour	7.1.x		
8.2	Middle point of outer lower lip contour	7.1.x		
8.3	Left corner of outer lip contour			
8.4	Right corner of outer lip contour			
8.5	Midpoint between f.p. 8.3 and 8.1 in outer upper lip contour	(8.3.x+8.1.x)/2		
8.6	Midpoint between f.p. 8.4 and 8.1 in outer upper lip contour	(8.4.x+8.1.x)/2		
8.7	Midpoint between f.p. 8.3 and 8.2 in outer lower lip contour	(8.3.x+8.2.x)/2		
8.8	Midpoint between f.p. 8.4 and 8.2 in outer lower lip contour	(8.4.x+8.2.x)/2		
8.9	Right hiph point of Cupid’s bow			
8.10	Left hiph point of Cupid’s bow			
9.1	Left nostril border			
9.2	Right nostril border			
9.3	Nose tip	7.1.x		
9.4	Bottom right edge of nose			
9.5	Bottom left edge of nose			
9.6	Right upper edge of nose bone			
9.7	Left upper edge of nose bone			
9.8	Top of the upper teeth	7.1.x		
9.9	Bottom of the lower teeth	7.1.x		

9.10	Bottom of the upper teeth	7.1.x		
9.11	Top of the lower teeth	7.1.x		
9.12	Middle lower edge of nose bone (or nose bump)	7.1.x	(9.6.y + 9.3.y)/2 or nose bump	
9.13	Left lower edge of nose bone		(9.6.y +9.3.y)/2	
9.14	Right lower edge of nose bone		(9.6.y +9.3.y)/2	
9.15	Bottom middle edge of nose	7.1.x		
10.1	Top of left ear			
10.2	Top of right ear			
10.3	Back of left ear		(10.1.y+10.5.y)/2	
10.4	Back of right ear		(10.2.y+10.6.y)/2	
10.5	Bottom of left ear lobe			
10.6	Bottom of right ear lobe			
10.7	Lower contact point between left lobe and face			
10.8	Lower contact point between right lobe and face			
10.9	Upper contact point between left ear and face			
10.10	Upper contact point between right ear and face			
11.1	Middle border between hair and forehead	7.1.x		
11.2	Right border between hair and forehead	< 4.4.x		
11.3	Left border between hair and forehead	> 4.3.x		
11.4	Top of skull	7.1.x		> 10.4.z and < 10.2.z
11.5	Hair thickness over f.p. 11.4	11.4.x		11.4.z
11.6	Back of skull	7.1.x	3.5.y	

Table C-4 -- FDP fields

FDP field	Description
featurePointsCoord	contains a **Coordinate** node. Specifies feature points for the calibration of the proprietary face. The coordinates are listed in the 'point' field in the **Coordinate** node in the prescribed order, that a feature point with a lower label is listed before a feature point with a higher label (e.g. feature point 3.14 before feature point 4.1).
textureCoords	contains a **Coordinate** node. Specifies texture coordinates for the

	feature points. The coordinates are listed in the **point** field in the **Coordinate** node in the prescribed order, that a feature point with a lower label is listed before a feature point with a higher label (e.g. feature point 3.14 before feature point 4.1).
textureType	contains a hint to the decoder on the type of texture image, in order to allow better interpolation of texture coordinates for the vertices that are not feature points. If **textureType** is 0, the decoder should assume that the texture image is obtained by cylindrical projection of the face. If **textureType** is 1, the decoder should assume that the texture image is obtained by orthographic projection of the face.
faceDefTables	contains **faceDefTables** nodes. The behavior of FAPs is defined in this field for the face in **faceSceneGraph**.
faceSceneGraph	contains a **Group** node. In case of option 1, this can be used to contain a texture image as explained above. In case of option 2, this is the grouping node for face model rendered in the compositor and has to contain the face model. In this case, the effect of Facial Animation Parameters is defined in the **faceDefTablesfield**.

Table C-5 -- Values for viseme_select

viseme_select	phonemes	example
0	none	na
1	p, b, m	put, bed, mill
2	f, v	far, voice
3	T,D	think, that
4	t, d	tip, doll
5	k, g	call, gas
6	tS, dZ, S	chair, join, she
7	s, z	sir, zeal
8	n, l	lot, not
9	r	red
10	A:	car
11	e	bed
12	I	tip
13	Q	top
14	U	book

The symbolic constants and variables used in the syntax diagrams are listed in the table below. Every group and BAP is assumed to have a unique unsigned integer value assignment.

TABLE:

Name	Definition	Description

NBAP_GROUP	Array[1..BAP_NUM_GROUPS]	Number of baps in each group.
BAPS_IN_GROUP	Array[1..BAP_NUM_GROUPS] [1..MAX_BAPS]	List of BAPs belonging to each group

Name	Value	Definition
MAX_BAPS	22	Maximum number of BAPs that belong to any group
BAP_NUM_GROUPS	24	Total number of BAP groups
NUM_BAPS	296	Total number of BAPs

The detailed BAPs and the BAP groups are given below:

BAP ID	BAP NAME	DESCRIPTION	Quant step size	Min/Max I-Frame quantized values	Min/Max P-Frame quantized values
1	sacroiliac_tilt	Forward-backward motion of the pelvis in the sagittal plane	64	-960/ +960	-600/ +600
2	sacroiliac_torsion	Rotation of the pelvis along the body vertical axis (defined by skeleton root)	64	-960/ +960	-600/ +600
3	sacroiliac_roll	Side to side swinging of the pelvis in the coronal plane	64	-960/ +960	-600/ +600
4	l_hip_flexion	Forward-backward rotation in the sagittal plane	128	-1260/ +1260	-600/ +600
5	r_hip_flexion	Forward-backward rotation in the sagittal plane	128	-1260/ +1260	-600/ +600
6	l_hip_abduct	Sideward opening in the coronal plane	128	-960/ +480	-600/ +600
7	r_hip_abduct	Sideward opening in the coronal plane	128	-960/ +480	-600/ +600
8	l_hip_twisting	Rotation along the thigh axis	256	-960/ +960	-360/ +360
9	r_hip_twisting	Rotation along the thigh axis	256	-960/ +960	-360/ +360
10	l_knee_flexion	Flexion-extension of the leg in the sagittal plane	128	-1500/ +180	-600/ +600
11	r_knee_flexion	Flexion-extension of the leg in the sagittal plane	128	-1500/ +180	-600/ +600
12	l_knee_twisting	Rotation along the shank axis.	256	-960/ +960	-360/ +360
13	r_knee_twisting	Rotation along the shank axis.	256	-960/ +960	-360/ +360
14	l_ankle_flexion	Flexion-extension of the foot in the sagittal plane	128	-780/ +780	-600/ +600
15	r_ankle_flexion	Flexion-extension of the foot in the sagittal plane	128	-780/ +780	-600/ +600
16	l_ankle_twisting	Rotation along the knee axis	256	-960/ +960	-360/ +360
17	r_ankle_twisting	Rotation along the knee axis	256	-960/ +960	-360/ +360
18	l_subtalar_flexion	Sideward orientation of the foot	256	-780/	-600/

				+780	+600
19	r_subtalar_flexion	Sideward orientation of the foot	256	-780/ +780	-600/ +600
20	l_midtarsal_twisting	Internal twisting of the foot (also called navicular joint in anatomy)	256	-180/ +180	-120/ +120
21	r_midtarsal_twisting	Internal twisting of the foot (also called navicular joint in anatomy)	256	-180/ +180	-120/ +120
22	l_metatarsal_flexion	Up and down rotation of the toe in the sagittal plane	256	-780/ +780	-600/ +600
23	r_metatarsal_flexion	Up and down rotation of the toe in the sagittal plane	256	-780/ +780	-600/ +600
24	l_sternoclavicular_abduct	Up and down motion in the coronal plane	128	-60/ +240	-120/ +120
25	r_sternoclavicular_abduct	Up and down motion in the coronal plane	128	-240/ +60	-120/ +120
26	l_sternoclavicular_rotate	Rotation in the transverse plane	128	-120/ +120	-60/ +60
27	r_sternoclavicular_rotate	Rotation in the transverse plane	128	-120/ +120	-60/ +60
28	l_acromioclavicular_abduct	Up and down motion in the coronal plane	128	-60/ +360	-120/ +120
29	r_acromioclavicular_abduct	Up and down motion in the coronal plane	128	-360/ +60	-120/ +120
30	l_acromioclavicular_rotate	Rotation in the transverse plane	128	-360/ +360	-120/ +120
31	r_acromioclavicular_rotate	Rotation in the transverse plane	128	-360/ +360	-120/ +120
32	l_shoulder_flexion	Forward-backward motion in the sagittal plane	64	-1080/ +1860	-600/ +600
33	r_shoulder_flexion	Forward-backward motion in the sagittal plane	64	-1080/ +1860	-600/ +600
34	l_shoulder_abduct	Sideward motion in the coronal plane	64	-240/ +1860	-600/ +600
35	r_shoulder_abduct	Sideward motion in the coronal plane	64	-1860/ +240	-600/ +600
36	l_shoulder_twisting	Rotation along the scapular axis	256	-960/ +960	-360/ +360
37	r_shoulder_twisting	Rotation along the scapular axis	256	-960/ +960	-360/ +360
38	l_elbow_flexion	Flexion-extension of the arm in the sagittal plane	64	-60/ +1560	-600/ +600
39	r_elbow_flexion	Flexion-extension of the arm in the sagittal plane	64	-60/ +1560	-600/ +600
40	l_elbow_twisting	Rotation of the forearm along the upper arm axis.	256	-960/ +960	-360/ +360
41	r_elbow_twisting	Rotation of the forearm along the upper arm axis.	256	-960/ +960	-360/ +360
42	l_wrist_flexion	Rotation of the hand in the coronal plane	128	-960/ +960	-600/ +600
43	r_wrist_flexion	Rotation of the hand in the coronal plane	128	-960/ +960	-600/ +600

44	l_wrist_pivot	Rotation of the hand in the sagittal planes	128	-660/ +660	-360/ +360
45	r_wrist_pivot	Rotation of the hand in the sagittal planes	128	-660/ +660	-360/ +360
46	l_wrist_twisting	Rotation of the hand along the forearm axis	256	-660/ +660	-360/ +360
47	r_wrist_twisting	Rotation of the hand along the forearm axis	256	-660/ +660	-360/ +360
48	skullbase_roll	Sideward motion of the skull along the frontal axis	128	-1860/ +1860	-600/ +600
49	skullbase_torsion	Twisting of the skull along the vertical axis	128	-1860/ +1860	-600/ +600
50	skullbase_tilt	Forward-backward motion in the sagittal plane along a lateral axis	128	-1860/ +1860	-600/ +600
51	vc1_roll	Sideward motion of vertebra C1	256	-240/ +240	-120/ +120
52	vc1_torsion	Twisting of vertebra C1	256	-240/ +240	-120/ +120
53	vc1_tilt	Forward-backward motion of vertebra C1 in the sagittal plane	256	-240/ +240	-120/ +120
54	vc2_roll	Sideward motion of vertebra C2	128	-660/ +660	-360/ +360
55	vc2_torsion	Twisting of vertebra C2	128	-660/ +660	-360/ +360
56	vc2_tilt	Forward-backward motion of vertebra C2 in the sagittal plane	128	-660/ +660	-360/ +360
57	vc3_roll	Sideward motion of vertebra C3	256	-240/ +240	-120/ +120
58	vc3_torsion	Twisting of vertebra C3	256	-240/ +240	-120/ +120
59	vc3_tilt	Forward-backward motion of vertebra C3 in the sagittal plane	256	-240/ +240	-120/ +120
60	vc4_roll	Sideward motion of vertebra C4	128	-660/ +660	-360/ +360
61	vc4_torsion	Twisting of vertebra C4	128	-660/ +660	-360/ +360
62	vc4_tilt	Forward-backward motion of vertebra C4 in the sagittal plane	128	-660/ +660	-360/ +360
63	vc5_roll	Sideward motion of vertebra C5	256	-240/ +240	-120/ +120
64	vc5_torsion	Twisting of vertebra C5	256	-240/ +240	-120/ +120
65	vc5_tilt	Forward-backward motion of vertebra C5 in the sagittal plane	256	-240/ +240	-120/ +120
66	vc6_roll	Sideward motion of vertebra C6	256	-240/ +240	-120/ +120
67	vc6_torsion	Twisting of vertebra C6	256	-240/ +240	-120/ +120
68	vc6_tilt	Forward-backward motion of vertebra C6 in the sagittal plane	256	-240/ +240	-120/ +120
69	vc7_roll	Sideward motion of vertebra C7	256	-240/	-120/

				+240	+120
70	vc7_torsion	Twisting of vertebra C7	256	-240/ +240	-120/ +120
71	vc7_tilt	Forward-backward motion of vertebra C7 in the sagittal plane	256	-240/ +240	-120/ +120
72	vt1_roll	Sideward motion of vertebra T1	128	-660/ +660	-360/ +360
73	vt1_torsion	Twisting of vertebra T1	128	-660/ +660	-360/ +360
74	vt1_tilt	Forward-backward motion of vertebra T1 in the sagittal plane	128	-660/ +660	-360/ +360
75	vt2_roll	Sideward motion of vertebra T2	256	-240/ +240	-120/ +120
76	vt2_torsion	Twisting of vertebra T2	256	-240/ +240	-120/ +120
77	vt2_tilt	Forward-backward motion of vertebra T2 in the sagittal plane	256	-240/ +240	-120/ +120
78	vt3_roll	Sideward motion of vertebra T3	256	-240/ +240	-120/ +120
79	vt3_torsion	Twisting of vertebra T3	256	-240/ +240	-120/ +120
80	vt3_tilt	Forward-backward motion of vertebra T3 in the sagittal plane	256	-240/ +240	-120/ +120
81	vt4_roll	Sideward motion of vertebra T4	256	-240/ +240	-120/ +120
82	vt4_torsion	Twisting of vertebra T4	256	-240/ +240	-120/ +120
83	vt4_tilt	Forward-backward motion of vertebra T4 in the sagittal plane	256	-240/ +240	-120/ +120
84	vt5_roll	Sideward motion of vertebra T5	256	-240/ +240	-120/ +120
85	vt5_torsion	Twisting of vertebra T5	256	-240/ +240	-120/ +120
86	vt5_tilt	Forward-backward motion of vertebra T5 in the sagittal plane	256	-240/ +240	-120/ +120
87	vt6_roll	Sideward motion of vertebra T6	128	-660/ +660	-360/ +360
88	vt6_torsion	Twisting of vertebra T6	128	-660/ +660	-360/ +360
89	vt6_tilt	Forward-backward motion of vertebra T6 in the sagittal plane	128	-660/ +660	-360/ +360
90	vt7_roll	Sideward motion of vertebra T7	256	-240/ +240	-120/ +120
91	vt7_torsion	Twisting of vertebra T7	256	-240/ +240	-120/ +120
92	vt7_tilt	Forward-backward motion of vertebra T7 in the sagittal plane	256	-240/ +240	-120/ +120
93	vt8_roll	Sideward motion of vertebra T8	256	-240/ +240	-120/ +120
94	vt8_torsion	Twisting of vertebra T8	256	-240/ +240	-120/ +120

95	vt8_tilt	Forward-backward motion of vertebra T8 in the sagittal plane	256	-240/ +240	-120/ +120
96	vt9_roll	Sideward motion of vertebra T9	256	-240/ +240	-120/ +120
97	vt9_torsion	Twisting of vertebra T9	256	-240/ +240	-120/ +120
98	vt9_tilt	Forward-backward motion of vertebra T9 in the sagittal plane	256	-240/ +240	-120/ +120
99	vt_10roll	Sideward motion of vertebra T10	128	-660/ +660	-360/ +360
100	vt10_torsion	Twisting of vertebra T10	128	-660/ +660	-360/ +360
101	vt10_tilt	Forward-backward motion of vertebra T10 in sagittal plane	128	-660/ +660	-360/ +360
102	vt11_roll	Sideward motion of vertebra T11	256	-240/ +240	-120/ +120
103	vt11_torsion	Twisting of vertebra T11	256	-240/ +240	-120/ +120
104	vt11_tilt	Forward-backward motion of vertebra T11 in sagittal plane	256	-240/ +240	-120/ +120
105	vt12_roll	Sideward motion of vertebra T12	256	-240/ +240	-120/ +120
106	vt12_torsion	Twisting of vertebra T12	256	-240/ +240	-120/ +120
107	vt12_tilt	Forward-backward motion of vertebra T12 in sagittal plane	256	-240/ +240	-120/ +120
108	vl1_roll	Sideward motion of vertebra L1	128	-660/ +660	-360/ +360
109	vl1_torsion	Twisting of vertebra L1	128	-660/ +660	-360/ +360
110	vl1_tilt	Forward-backward motion of vertebra L1 in sagittal plane	128	-660/ +660	-360/ +360
111	vl2_roll	Sideward motion of vertebra L2	256	-240/ +240	-120/ +120
112	vl2_torsion	Twisting of vertebra L2	256	-240/ +240	-120/ +120
113	vl2_tilt	Forward-backward motion of vertebra L2 in sagittal plane	256	-240/ +240	-120/ +120
114	vl3_roll	Sideward motion of vertebra L3	128	-660/ +660	-360/ +360
115	vl3_torsion	Twisting of vertebra L3	128	-660/ +660	-360/ +360
116	vl3_tilt	Forward-backward motion of vertebra L3 in sagittal plane	128	-660/ +660	-360/ +360
117	vl4_roll	Sideward motion of vertebra L4	256	-240/ +240	-120/ +120
118	vl4_torsion	Twisting of vertebra L4	256	-240/ +240	-120/ +120
119	vl4_tilt	Forward-backward motion of vertebra L4 in sagittal plane	256	-240/ +240	-120/ +120
120	vl5_roll	Sideward motion of vertebra L5	128	-660/	-360/

				+660	+360
121	vl5_torsion	Twisting of vertebra L5	128	-660/ +660	-360/ +360
122	vl5_tilt	Forward-backward motion of vertebra L5 in sagittal plane	128	-660/ +660	-360/ +360
123	l_pinky0_flexion	Metacarpal flexing mobility of the pinky finger	512	-240/ +240	-120/ +120
124	r_pinky0_flexion	Metacarpal flexing mobility of the pinky finger	512	-240/ +240	-120/ +120
125	l_pinky1_flexion	First knukle of the pinky finger	128	-1140/ +240	-240/ +240
126	r_pinky1_flexion	First knukle of the pinky finger	128	-1140/ +240	-240/ +240
127	l_pinky1_pivot	Lateral mobility of the pinky finger	128	-420/ +120	-120/ +120
128	r_pinky1_pivot	Lateral mobility of the pinky finger	128	-120/ +420	-120/ +120
129	l_pinky1_twisting	Along the pinky finger axis	256	-180/ +360	-120/ +120
130	r_pinky1_twisting	Along the pinky finger axis	256	-360/ +180	-120/ +120
131	l_pinky2_flexion	Second knuckle of the pinky number	128	-1380/ +60	-240/ +240
132	r_pinky2_flexion	Second knuckle of the pinky number	128	-1380/ +60	-240/ +240
133	l_pinky3_flexion	Third knuckle of the pinky finger	128	-960/ +60	-240/ +240
134	r_pinky3_flexion	Third knuckle of the pinky finger	128	-960/ +60	-240/ +240
135	l_ring0_flexion	Metacarpal flexing mobility of the ring finger	256	-180/ +180	-120/ +120
136	r_ring0_flexion	Metacarpal flexing mobility of the ring finger	256	-180/ +180	-120/ +120
137	l_ring1_flexion	First knukle of the ring finger	128	-1140/ +360	-240/ +240
138	r_ring1_flexion	First knukle of the ring finger	128	-1140/ +360	-240/ +240
139	l_ring1_pivot	Lateral mobility of the ring finger	128	-240/ +120	-120/ +120
140	r_ring1_pivot	Lateral mobility of the ring finger	128	-120/ +240	-120/ +120
141	l_ring1_twisting	Along the ring finger axis	256	-240/ +240	-120/ +120
142	r_ring1_twisting	Along the ring finger axis	256	-240/ +240	-120/ +120
143	l_ring2_flexion	Second knuckle of the ring number	128	-1380/ +60	-240/ +240
144	r_ring2_flexion	Second knuckle of the ring number	128	-1380/ +60	-240/ +240
145	l_ring3_flexion	Third knuckle of the ring finger	128	-960/	-240/

				+60	+240
146	r_ring3_flexion	Third knuckle of the ring finger	128	-960/ +60	-240/ +240
147	l_middle0_flexion	Metacarpal flexing mobility of the middle finger	256	-120/ +120	-120/ +120
148	r_middle0_flexion	Metacarpal flexing mobility of the middle finger	256	-120/ +120	-120/ +120
149	l_middle1_flexion	First knukle of the middle finger	128	-1140/ +360	-240/ +240
150	r_middle1_flexion	First knukle of the middle finger	128	-1140/ +360	-240/ +240
151	l_middle1_pivot	Lateral mobility of the middle finger	128	-180/ +180	-120/ +120
152	r_middle1_pivot	Lateral mobility of the middle finger	128	-180/ +180	-120/ +120
153	l_middle1_twisting	Along the middle finger axis	256	-180/ +180	-120/ +120
154	r_middle1_twisting	Along the middle finger axis	256	-180/ +180	-120/ +120
155	l_middle2_flexion	Second knuckle of the middle number	128	-1380/ +60	-240/ +240
156	r_middle2_flexion	Second knuckle of the middle number	128	-1380/ +60	-240/ +240
157	l_middle3_flexion	Third knuckle of the middle finger	128	-960/ +60	-240/ +240
158	r_middle3_flexion	Third knuckle of the middle finger	128	-960/ +60	-240/ +240
159	l_index0_flexion	Metacarpal flexing mobility of the index finger	256	-60/ +60	-60/ +60
160	r_index0_flexion	Metacarpal flexing mobility of the index finger	256	-60/ +60	-60/ +60
161	l_index1_flexion	First knukle of the index finger	128	-1140/ +360	-240/ +240
162	r_index1_flexion	First knukle of the index finger	128	-1140/ +360	-240/ +240
163	l_index1_pivot	Lateral mobility of the index finger	128	-120/ +240	-60/ +60
164	r_index1_pivot	Lateral mobility of the index finger	128	-240/ +120	-60/ +60
165	l_index1_twisting	Along the index finger axis	256	-240/ +180	-120/ +120
166	r_index1_twisting	Along the index finger axis	256	-180/ +240	-120/ +120
167	l_index2_flexion	Second knuckle of the index number	128	-1380/ +60	-240/ +240
168	r_index2_flexion	Second knuckle of the index number	128	-1380/ +60	-240/ +240
169	l_index3_flexion	Third knuckle of the index finger	128	-960/ +60	-240/ +240
170	r_index3_flexion	Third knuckle of the index finger	128	-960/	-240/

				+60	+240
171	l_thumb1_flexion	First knukle of the thumb finger	128	-480/ +960	-240/ +240
172	r_thumb1_flexion	First knukle of the thumb finger	128	-960/ +480	-240/ +240
173	l_thumb1_pivot	Lateral mobility of the thumb finger	128	-120/ +1080	-240/ +240
174	r_thumb1_pivot	Lateral mobility of the thumb finger	128	-1080/ +120	-240/ +240
175	l_thumb1_twisting	Along the thumb finger axis	256	-720/ +120	-240/ +240
176	r_thumb1_twisting	Along the thumb finger axis	256	-720/ +120	-240/ +240
177	l_thumb2_flexion	Second knuckle of the thumb number	128	-780/ +60	-240/ +240
178	r_thumb2_flexion	Second knuckle of the thumb number	128	-780/ +60	-240/ +240
179	l_thumb3_flexion	Third knuckle of the thumb finger	128	-960/ +60	-240/ +240
180	r_thumb3_flexion	Third knuckle of the thumb finger	128	-960/ +60	-240/ +240
181	HumanoidRoot_tr_vertical	Body origin translation in vertical direction	64	-1860/ +1860	-600/ +600
182	HumanoidRoot_tr_lateral	Body origin translation in lateral direction	64	-1860/ +1860	-600/ +600
183	HumanoidRoot_tr_frontal	Body origin translation in frontal direction	64	-1860/ +1860	-600/ +600
184	HumanoidRoot_rt_body_turn	Rotation of the skeleton root along the body coordinate system's vertical axis	64	-1860/ +1860	-600/ +600
185	HumanoidRoot_rt_body_roll	Rotation of the skeleton root along the body coordinate system's frontal axis	64	-1860/ +1860	-600/ +600
186	HumanoidRoot_rt_body_tilt	Rotation of the skeleton root along the body coordinate system's lateral axis	64	-1860/ +1860	-600/ +600

BAPs 187 through 296 are extension BAPs for use with the BodyDefTable nodes., see Systems CD for specification of BodyDefTable nodes. These BAPs are user-defined. The step sizes of these BAPs are 1, and I-Frame and P-Frame default minmax values are not defined.

GROUP ID	**GROUP NAME**	**BAPS**
	Pelvis	sacroiliac_tilt, sacroiliac_torsion, sacroiliac_roll (1,2,3)
	Left leg1	l_hip_flexion, l_hip_abduct, l_knee_flexion, l_ankle_flexion (4,6,10,14)
	Right leg1	r_hip_flexion, r_hip_abduct, r_knee_flexion, r_ankle_flexion (5,7,11,15)
	Left leg2	l_hip_twisting, l_knee_twisting, l_ankle_twisting, l_subtalar_flexion, l_midtarsal_flexion, l_metatarsal_flexion (8,12,16,18,20,22)
	Right leg2	r_hip_twisting, r_knee_twisting, r_ankle_twisting, r_subtalar_flexion, r_midtarsal_flexion, r_metatarsal_flexion (9,13,17,19,21,23)
	Left arm1	l_shoulder_flexion, l_shoulder_abduct, l_shoulder_twisting, l_elbow_flexion, l_wrist_flexion

		(32,34,36,38,42)
	Right arm1	r_shoulder_flexion, r_shoulder_abduct, r_shoulder_twisting, r_elbow_flexion, r_wrist_flexion (33,35,37,39,43)
	Left arm2	l_sternoclavicular_abduct, l_sternoclavicular_rotate, l_acromioclavicular_abduct, l_acromioclavicular_rotate, l_elbow_twisting, l_wrist_pivot, l_wrist_twisting (24,26,28,30,40,44,46)
	Right arm2	r_sternoclavicular_abduct, r_sternoclavicular_rotate, r_acromioclavicular_abduct, r_acromioclavicular_rotate, r_elbow_twisting, r_wrist_pivot, r_wrist_twisting (25,27,29,31,41,45,47)
	Spine1	skullbase_roll, skullbase_torsion, skullbase_tilt, vc4roll, vc4torsion, vc4tilt, vt6roll, vt6torsion, vt6tilt, vl3roll, vl3torsion, vl3tilt, (48,49,50,60,61,62,87,88,89,114,115,116)
	Spine2	vc2roll, vc2torsion, vc2tilt, vt1roll, vt1torsion, vt1tilt, vt10roll, vt10torsion, vt10tilt, vl1roll, vl1torsion, vl1tilt, vl5roll, vl5torsion, vl5tilt (54,55,56,72,73,74,99,100,101,108, 109,110,120,121,122)
	Spine3	vc3roll, vc3torsion, vc3tilt, vc6roll, vc6torsion, vc6tilt, vt4roll, vt4torsion, vt4tilt, vt8roll, vt8torsion, vt8tilt, vt12roll, vt12torsion, vt12tilt vl4roll, vl4torsion, vl4tilt, (57,58,59,66,67,68,81,82,83,93,94,95, 105,106,107,117,118,119)
	Spine4	vc5roll, vc5torsion, vc5tilt, vc7roll, vc7torsion, vc7tilt vt2roll, vt2torsion, vt2tilt, vt7roll, vt7torsion, vt7tilt, vt11roll, vt11torsion, vt11tilt, vl2roll, vl2torsion, vl2tilt, (63,64,65,69,70,71,75,76,77,90,91, 92,102,103,104,111,112,113)
	Spine5	vc1roll, vc1torsion, vc1tilt, vt3roll, vt3torsion, vt3tilt, vt5roll, vt5torsion, vt5tilt, vt9roll, vt9torsion, vt9tilt, (51,52,53,78,79,80,84,85,86,96,97,98)
	Left hand1	l_pinky1_flexion, l_pinky2_flexion, l_pinky3_flexion, l_ring1_flexion, l_ring2_flexion, l_ring3_flexion, l_middle1_flexion, l_middle2_flexion,l_middle3_flexion, l_index1_flexion, l_index2_flexion, l_index3_flexion, l_thumb1_flexion, l_thumb1_pivot, l_thumb2_flexion, l_thumb3_flexion (125,131,133,137,143,145,149,155,157, 161,167,169,171,173,177,179)
	Right hand1	r_pinky1_flexion, r_pinky2_flexion, r_pinky3_flexion, r_ring1_flexion, r_ring2_flexion, r_ring3_flexion,

		r_middle1_flexion, r_middle2_flexion,r_middle3_flexion, r_index1_flexion, r_index2_flexion, r_index3_flexion, r_thumb1_flexion, r_thumb1_pivot, r_thumb2_flexion, r_thumb3_flexion (126,132,134,138,144,146,150,156,158, 162,168,170,172,174,178,180)
	Left hand2	l_pinky0_flexion, l_pinky1_pivot, l_pinky1_twisting, l_ring0_flexion, l_ring1_pivot, l_ring1_twisting, l_middle0_flexion, l_middle1_pivot, l_middle1_twisting, l_index0_flexion, l_index1_pivot, l_index1_twisting, l_thumb1_twisting (123,127,129,135,139,141,147, 151,153,159,163,165,175)
	Right hand2	r_pinky0_flexion, r_pinky1_pivot, r_pinky1_twisting, r_ring0_flexion, r_ring1_pivot, r_ring1_twisting, r_middle0_flexion, r_middle1_pivot, r_middle1_twisting, r_index0_flexion, r_index1_pivot, r_index1_twisting, r_thumb1_twisting (124,128,130,136,140,142,148, 152,154,160,164,166,176)
	Global positioning	HumanoidRoot_tr_vertical, HumanoidRoot_tr_lateral, HumanoidRoot_tr_frontal, HumanoidRoot_rt_body_turn, HumanoidRoot_rt_body_roll, HumanoidRoot_rt_body_tilt (181,182,183,184,185,186)

Additionally, groups 20 to 24 contain extension BAPs. Extension BAPs are user-defined BAPs for use with BodyDefTables. See Systems CD for specification of BodyDefTable nodes. Groups 20 to 24 each contain 22 BAPs. Thus, the extension BAP groups are as follows:

GROUP ID	**GROUP NAME**	**BAPs**
20	Extension1	187…208
21	Extension2	209…230
22	Extension3	231…252
23	Extension4	253…274
24	Extension5	275…296

Detailed Degrees of Freedom

The ***degrees of freedom (DOF)*** sufficient to locate and animate a human body are decomposed into six DOF for the global location and 180 DOF for the internal mobilities. The topology of the suggested joints is given below. The hands are optional with an additional set of joints.

Suggested default joint center values for broadcast applications

HumanoidRoot	0.0000	0.9723	-0.0728
sacroiliac	0.0000	0.9710	-0.0728
l_hip	0.0956	0.9364	0.0000
l_knee	0.0956	0.5095	-0.0036
l_ankle	0.0946	0.0762	-0.0261
l_subtalar	0.0956	0.0398	0.0069
l_midtarsal	0.1079	0.0317	0.0670
l_metatarsal	0.0942	0.0092	0.1239

r_hip	-0.0956	0.9364	0.0000
r_knee	-0.0956	0.5095	-0.0036
r_ankle	-0.0946	0.0762	-0.0261
r_subtalar	-0.0956	0.0398	0.0069
r_midtarsal	-0.1079	0.0317	0.0670
r_metatarsal	-0.0942	0.0092	0.1239
vl5	0.0000	1.0817	-0.0728
vl4	0.0000	1.1174	-0.0727
vl3	0.0000	1.1525	-0.0727
vl2	0.0000	1.1795	-0.0727
vl1	0.0000	1.2161	-0.0727
vt12	0.0000	1.2527	-0.0727
vt11	0.0000	1.2918	-0.0727
vt10	0.0000	1.3098	-0.0737
vt9	0.0000	1.3375	-0.0752
vt8	0.0000	1.3631	-0.0758
vt7	0.0000	1.3875	-0.0745
vt6	0.0000	1.4116	-0.0712
vt5	0.0000	1.4351	-0.0657
vt4	0.0000	1.4569	-0.0587
vt3	0.0000	1.4832	-0.0482
vt2	0.0000	1.5011	-0.0397
vt1	0.0000	1.5201	-0.0300
vc7	0.0000	1.5382	-0.0213
vc6	0.0000	1.5607	-0.0073
vc5	0.0000	1.5770	-0.0012
vc4	0.0000	1.5912	-0.0014
vc3	0.0000	1.6050	-0.0033
vc2	0.0000	1.6178	-0.0033
vc1	0.0000	1.6394	0.0036
skullbase	0.0000	1.6440	0.0036
l_sternoclavicular	0.0757	1.4844	-0.0251
l_acromioclavicular	0.0899	1.4525	-0.0322
l_shoulder	0.1968	1.4642	-0.0265
l_elbow	0.1982	1.1622	-0.0557
l_wrist	0.1972	0.8929	-0.0690
l_thumb1	0.1912	0.8734	-0.0657
l_thumb2	0.1912	0.8156	-0.0079
l_thumb3	0.1912	0.8007	0.0070
l_index0	0.1912	0.8259	-0.0460

l_index1	0.1912	0.8050	-0.0460
l_index2	0.1912	0.7595	-0.0460
l_index3	0.1912	0.7370	-0.0460
l_middle0	0.1912	0.8282	-0.0710
l_middle1	0.1912	0.8071	-0.0710
l_middle2	0.1912	0.7525	-0.0710
l_middle3	0.1912	0.7263	-0.0710
l_ring0	0.1912	0.8302	-0.0972
l_ring1	0.1912	0.8098	-0.0972
l_ring2	0.1912	0.7570	-0.0972
l_ring3	0.1912	0.7328	-0.0972
l_pinky0	0.1912	0.8373	-0.1211
l_pinky1	0.1912	0.8173	-0.1211
l_pinky2	0.1912	0.7759	-0.1211
l_pinky3	0.1912	0.7584	-0.1211
r_sternoclavicular	-0.0757	1.4844	-0.0251
r_acromioclavicular	-0.0899	1.4525	-0.0322
r_shoulder	-0.1968	1.4642	-0.0265
r_elbow	-0.1982	1.1622	-0.0557
r_wrist	-0.1972	0.8929	-0.0690
r_thumb1	-0.1912	0.8734	-0.0657
r_thumb2	-0.1912	0.8156	-0.0079
r_thumb3	-0.1912	0.8007	0.0070
r_index0	-0.1912	0.8259	-0.0460
r_index1	-0.1912	0.8050	-0.0460
r_index2	-0.1912	0.7595	-0.0460
r_index3	-0.1912	0.7370	-0.0460
r_middle0	-0.1912	0.8282	-0.0710
r_middle1	-0.1912	0.8071	-0.0710
r_middle2	-0.1912	0.7525	-0.0710
r_middle3	-0.1912	0.7263	-0.0710
r_ring0	-0.1912	0.8302	-0.0972
r_ring1	-0.1912	0.8098	-0.0972
r_ring2	-0.1912	0.7570	-0.0972
r_ring3	-0.1912	0.7328	-0.0972
r_pinky0	-0.1912	0.8373	-0.1211
r_pinky1	-0.1912	0.8173	-0.1211
r_pinky2	-0.1912	0.7759	-0.1211
r_pinky3	-0.1912	0.7584	-0.1211

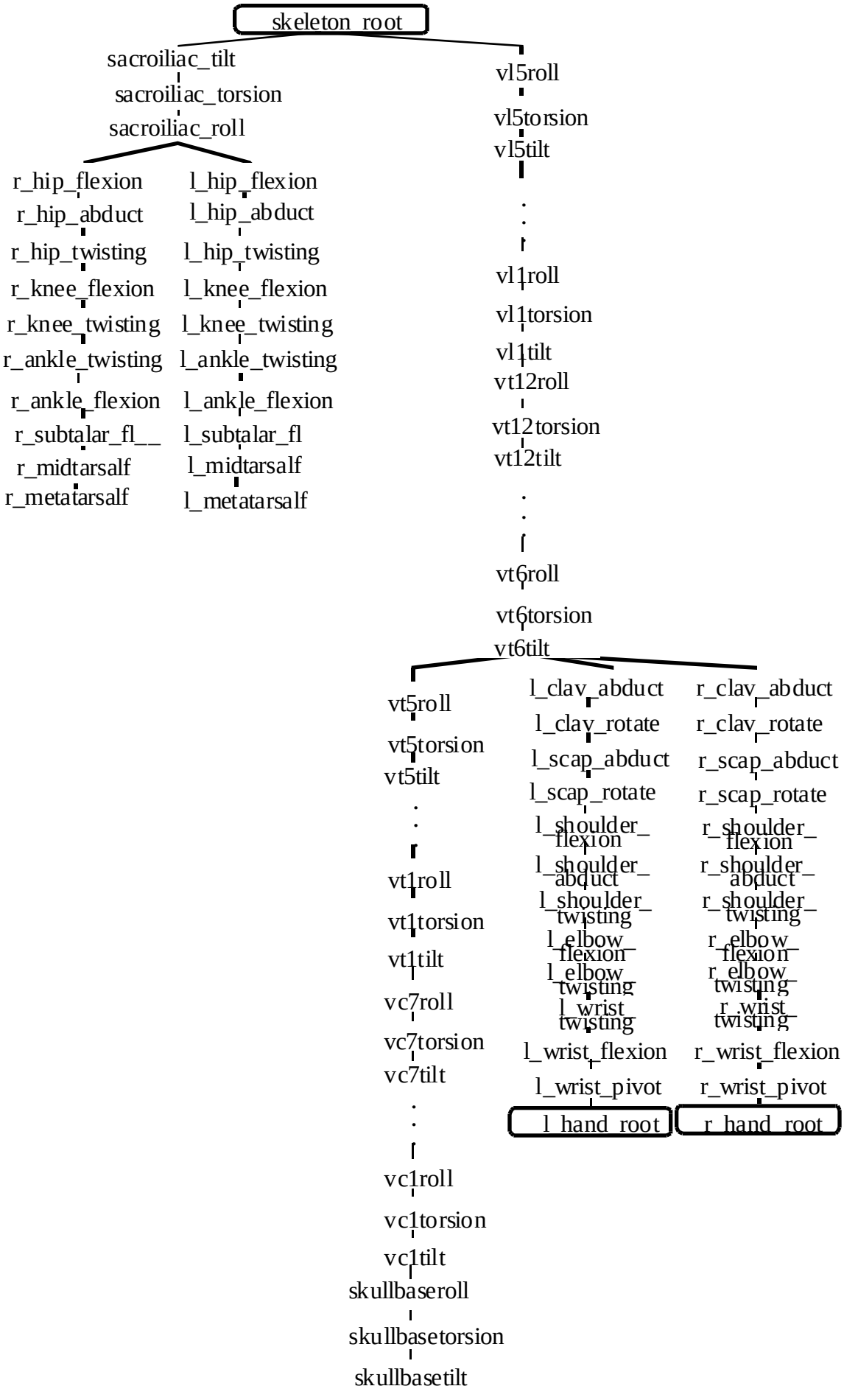

skeleton_root
sacroiliac_tilt
sacroiliac_torsion
sacroiliac_roll
r_hip_flexion
r_hip_abduct
r_hip_twisting
r_knee_flexion
r_knee_twisting
r_ankle_twisting
r_ankle_flexion
r_subtalar_fl__
r_midtarsalf
r_metatarsalf
l_hip_flexion
l_hip_abduct
l_hip_twisting
l_knee_flexion
l_knee_twisting
l_ankle_twisting
l_ankle_flexion
l_subtalar_fl
l_midtarsalf
l_metatarsalf
vl5roll
vl5torsion
vl5tilt
vl1roll
vl1torsion
vl1tilt
vt12roll
vt12torsion
vt12tilt
vt6roll
vt6torsion
vt6tilt
vt5roll
vt5torsion
vt5tilt
vt1roll
vt1torsion
vt1tilt
vc7roll
vc7torsion
vc7tilt
vc1roll
vc1torsion
vc1tilt
skullbaseroll
skullbasetorsion
skullbasetilt
l_clav_abduct
l_clav_rotate
l_scap_abduct
l_scap_rotate
l_shoulder_flexion
l_shoulder_abduct
l_shoulder_twisting
l_elbow_flexion
l_elbow_twisting
l_wrist_twisting
l_wrist_flexion
l_wrist_pivot
l_hand_root
r_clav_abduct
r_clav_rotate
r_scap_abduct
r_scap_rotate
r_shoulder_flexion
r_shoulder_abduct
r_shoulder_twisting
r_elbow_flexion
r_elbow_twisting
r_wrist_twisting
r_wrist_flexion
r_wrist_pivot
r_hand_root

Figure V2 -4 -- Body topology.

Lower Body

From the skeleton_root node, three DOF allow flexible pelvic orientation followed by nine DOF per leg, from the hip joint to the toe joint.

First, the pelvic mobilities are very important to convey a gender personification to the motion. The naming convention of these mobilities is also used for the spine DOF. The degrees of freedom are:

sacroiliac_tilt : forward-backward motion in the sagital plane

sacroiliac_torsion :rotation along the body_vertical axis (defined by_skeleton root)

sacroiliac_roll : side to side swinging in the coronal plane

The leg mobilities follow in this order :

At the hip :

hip_flexion : forward-backward rotation in the sagital plane

hip_abduct : sideward opening in the coronal plane

hip_twisting : rotation along the thigh axis.

At the knee :

knee_flexion : flexion-extension of the leg in the sagital plane

knee_twisting : rotation along the shank axis.

At the ankle :

ankle_twisting : rotation along the shank axis. This joint is redundant with the knee_twisting except that only the foot rotate and not the shank segment.

ankle_flexion : flexion-extension of the foot_in the sagital plane

At the foot complex:

The foot complex region is completely described with three degrees of freedom with independent position and orientation : the subtalar joint, the *mid_foot* joint, between the subtalar joint and the toe joint to capture complex internal relative movement of the foot bones (also called the navicular joint in the literature).

subtalar_flexion : sideward orientation of the foot

midtarsal_flexion : internal twisting of the foot (also called navicular joint in anatomy)

metatorsal_flexion : up and down rotation of the toe in the sagital plane

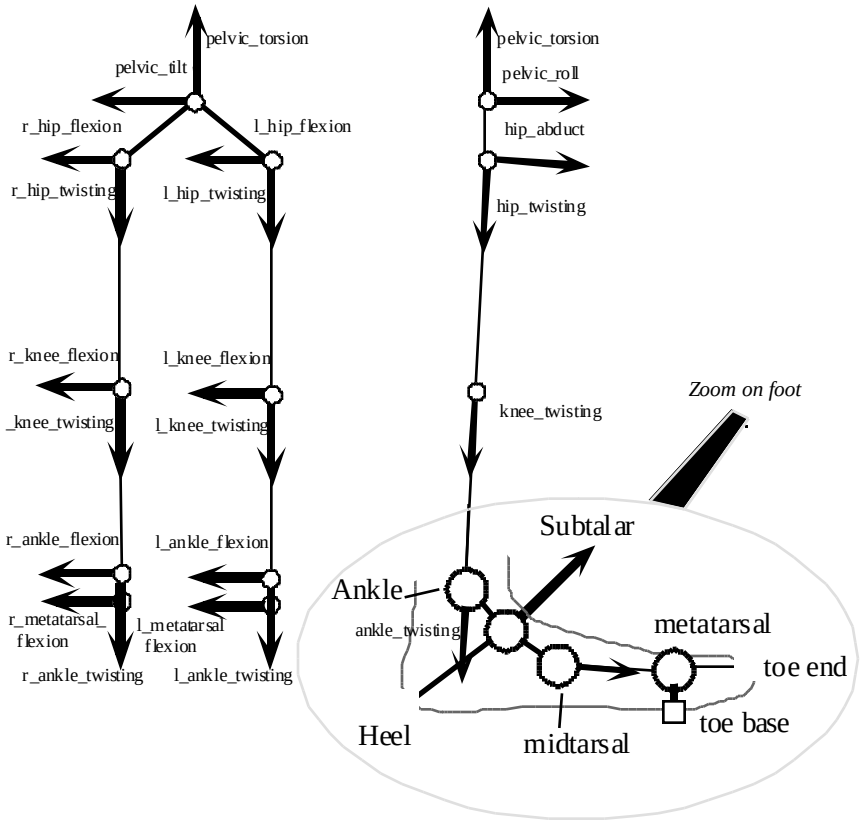


Figure V2 -5 -- Front and side views of the mobilities of the leg

Upper Body

The suggested upper body degrees of freedom and their corresponding axes of rotation are shown in the following diagrams.

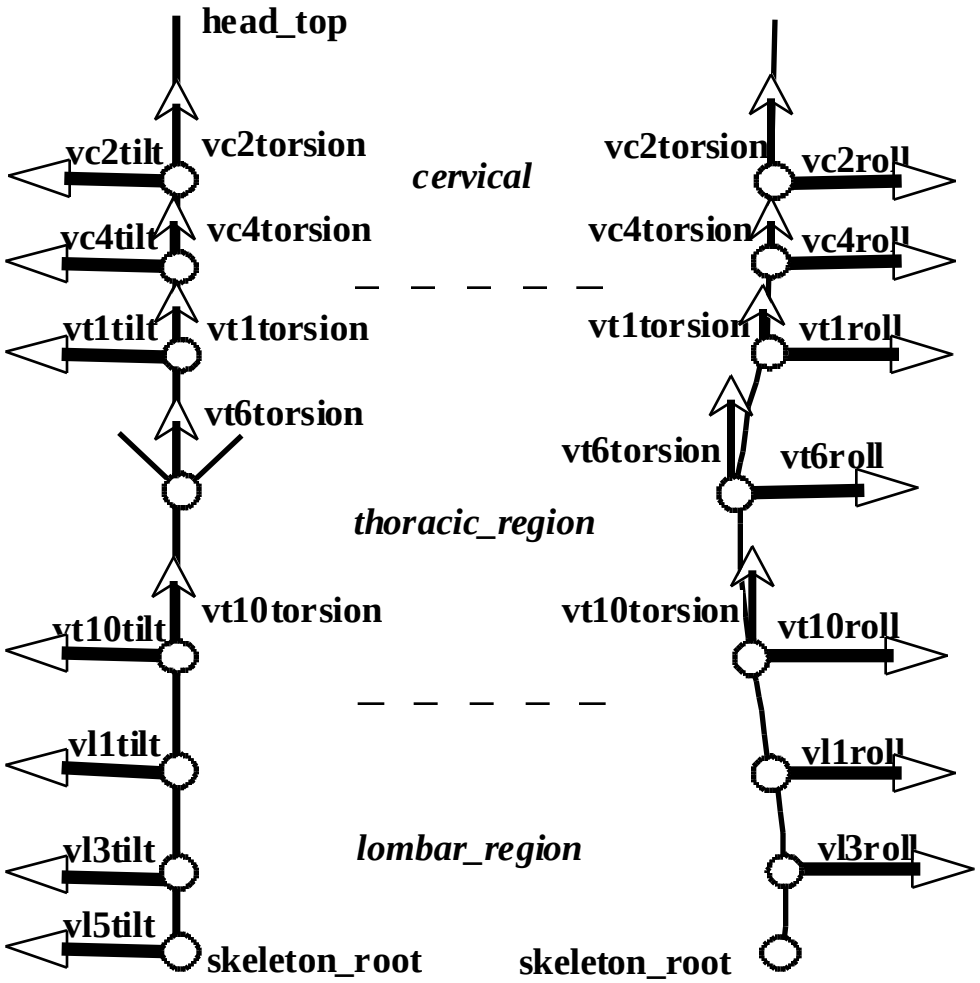


Figure V2 -6 -- Front and side views of the mobilities of the simple spine

At the spine:
The vertebrae are dispatched into 5 *lumbar*, 12 *thoracic* and *cervical* groups.A total of 72 DOFs are defined for the spine, therefore complex applications with whole spine mobilities can be developed. However, typically, the application will use simpler spines that balance the computational speed, with the realism of animation. Therefore, 5 groups of spine are defined, from the simplest to most complex. It is suggested that the spine groups (Spine1, Spine2) are used for simple spine.

At the arms:
The arm structure is attached to the spine. The mobilities are similar to the ones defined for the leg when the arm is twisted such that the hand palm is facing backward. The name of the DOF respects this structure similarity.
Strictly speaking, the two clavicula DOF and the two scapula DOF are not part of the arm structure, they only initiate its articulated chain. The scapula joints improve the mechanical model of the shoulder complex. It should be noted that such a chain representation is only a step toward a mechanically correct representation of this region, mainly because the shoulder complex is in fact a closed loop mechanism. The scapula holds the same mobility as the clavicula but very close to the shoulder joint.

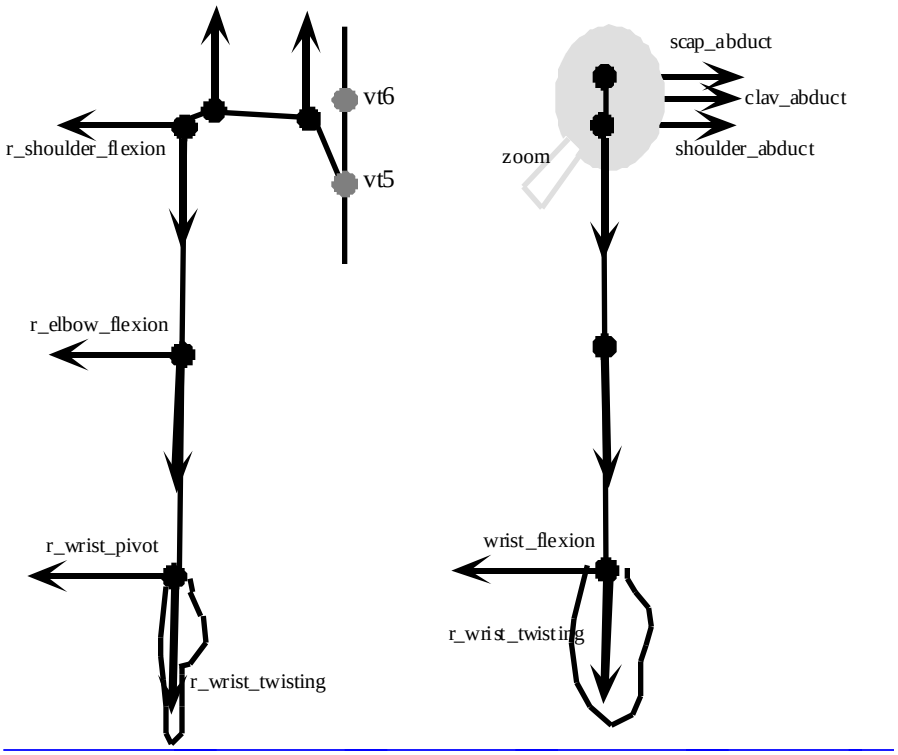


Figure V2 -7 -- Front and side views of the mobilities of the arm in rest position

At the clavicula :
- **sternoclavicular_abduct** : up and down motion in the coronal plane
- **sternoclavicular_rotate** : rotation in the transverse plane

At the scapula :
- **acromioclavicular_abduct** : up and down motion in the coronal plane
- **acromioclavicular_rotate** : rotation in the transverse plane

At the shoulder :
- **shoulder_flexion** : forward-backward motion in the sagital plane
- **shoulder_abduct** : sideward motion in the coronal plane
- **shoulder_twisting** : along the forearm axis

At the elbow :
- **elbow_flexion** : flexion-extension of the arm in the sagital plane
- **elbow_twisting** : along the arm axis.

At the wrist :
- **wrist_twisting** : along the arm axis. This DOF is redundant with the elbow twisting except that only the hand rotate and not the forearm
- **wrist_flexion** : rotation of the hand in the coronal plane
- **wrist_pivot** : rotation of the hand in the sagital planes

Head rotations

Note that there are three BAPs defined for head rotation : skullbase_roll, skullbase_torsion, skullbase_tilt . There are also 3 FAPs for head rotation. If both FAPs and BAPs are used for head rotation, then these FAPs shall denote the head rotation with respect to the skullbase coordinate system.

Hands

The hand mobilities have a standard structure for the five fingers. This structure is organized as :
- a **flexing** DOF for closing the first knuckle
- a **pivoting** rotation for the lateral mobility of the finger
- a **twisting** rotation for small adjustments of the finger grasping orientation

two other **flexing** DOF for closing the second and third knuckles

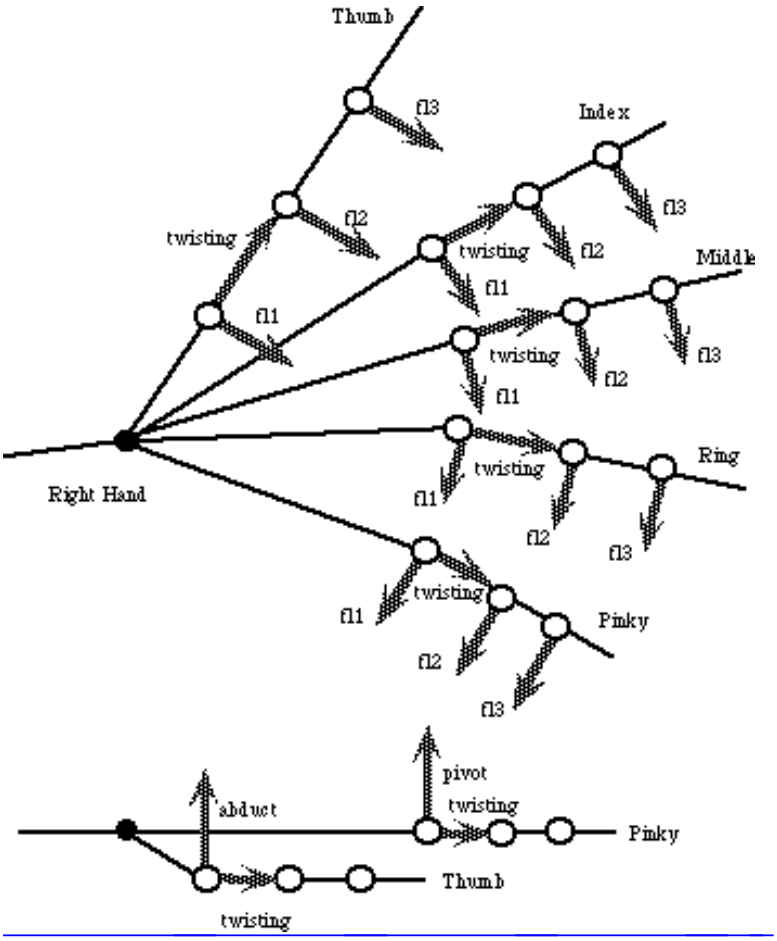


Figure V2 -8 -- Mobilities of the right hand

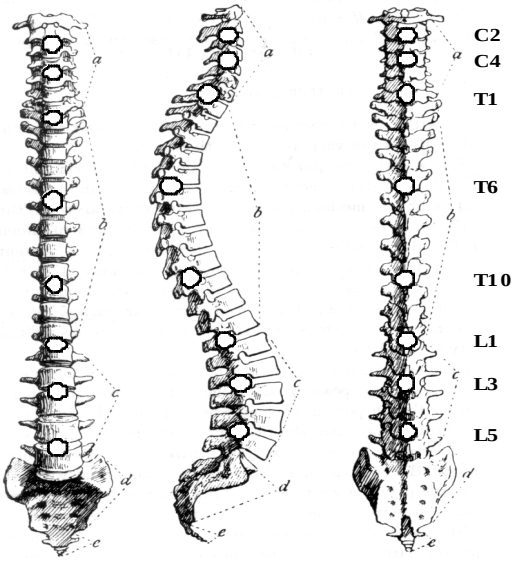


Figure V2 -9 -- Example complex spine mobilities and their simplified model names.

Integration of Facial Animation with TTS

The following figure shows the complete block diagram describing the integration of a proprietary TTS Synthesizer into an ISO/IEC 14496 face animation system. The FAP bookmarks defined by the user in the input text of the TTS Stream are identified by the speech synthesizer and transmitted in ASCII format to the Phoneme to FAP converter using the TtsFAPInterface as defined in ISO/IEC 14496-3.

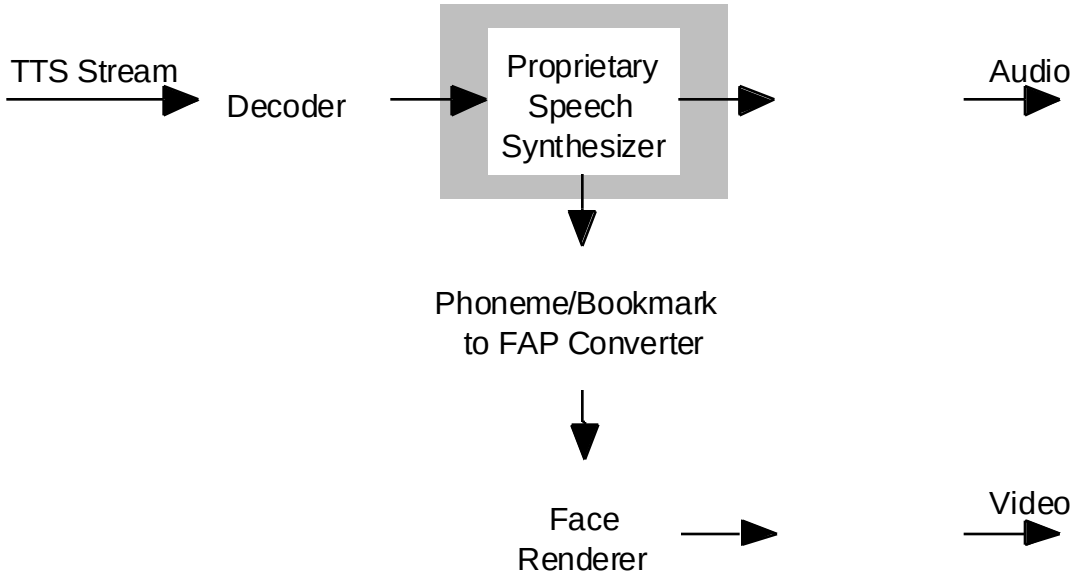


Figure C-2: Blockdiagram showing the integration of a proprietary TTS into an ISO/IEC 14496 face animation system with Phoneme/Bookmark to FAP Converter.

The syntax of the bookmark sequences used to convey commands to the TTS system is the following:

<FAP *n* FAPfields *T C* >

n: FAP number defined according to annex C, Table C-1 with 2<=n<=68.

FAPfields := expression_select1* expression_intensity1 expression_select2* expression_intensity2, in case n == 2

FAPfields := *a***, in case 2 < n < =68

T: transition time defined in ms

C: time curve for computation of *a* during transition time *T*

where ":=" indicates a production rule
* defined according to Table C-3 (expression select)

** defined in units according to Table C-1

Phoneme/Bookmark to FAP Converter

In addition to its function to convert phonemes into visemes, the Phoneme/Bookmark to FAP Converter (Fig. 1) is responsible for translating the FAP bookmarks defined by the user and made available by the bookmark field of the TtsFAPInterface as defined in ISO/IEC 14496-1 into a sequence of FAPs that can be interpreted by the Face Renderer, which then has to use them. An interpolation function is used for computing the FAP amplitudes. The converter merges the FAPs it derives from phonemes and bookmarks into one set of FAP parameters for each time instant using the following steps:

1. Set all low-level FAPs to interpolate.
2. Set specified FAPs to values computed using the interpolation function and phonemes.

3. init_face is set according to the following suggestion: In case of FAP 1 viseme_select1/2 != 0 and FAP 2 expression_select1/2 !=0, init_face=1; In case of FAP 2 expression_select1/2 !=0 and using FAP 3-68 for visemes, init_face=0.

Facial expressions and visemes are combined such that mouth closures are achieved for the relevant visemes while maintaining the overall facial expression. In case that the bookmarks contain visemes, they are ignored.

Transitions between facial expressions are achieved by computing the interpolation function for expression_intensity1 and expression_intensity2. Smooth transitions have to be achieved if expression_select1 and expression_select2 are the same for consecutive bookmarks.

Face Renderer:

In the case that the face model is also animated from an FAP parameter stream, the face renderer has to animate the face using the input from the TtsFAPInterface which is processed by the Phoneme/Bookmark to FAP converter and the input from the FAP decoder. In case that the renderer receives values for the same FAP from both sources, the values derived from the FAP stream take precedence.

Interpolation Functions

For simplicity, the following description on how to compute the amplitude of an FAP is explained using the amplitude *a*. The same method is applied to expression_intensity1 and expression_intensity2.

The FAP amplitude *a* defines the amplitude to be applied at the end of the transition time *T*. The amplitude a_s of the FAP at the beginning of the transition depends on previous bookmarks and can be equal to:

- 0 if the FAP bookmark is the first one with this FAP *n* made available through the TtsFAPInterface.
- *a* of the previous FAP bookmark with the same FAP *n* if a time longer than the previous transition time *T* has elapsed between these two FAP bookmarks.
- The actual reached amplitude due to the previous FAP definition if a time shorter than the previous transition time *T* has elapsed between the two FAP bookmarks.

At the end of the transition time *T*, *a* is maintained until another FAP bookmark gives a new value to reach. To reset an FAP, a bookmark for FAP *n* with *a*=0 is transmitted in the text.

To avoid too many parameters for defining the evolution of the amplitude during the transition time, the functions that compute for each frame the amplitude of the FAP to be sent to the face renderer are predefined. Assuming that the transition time *T* is always 1, the following 3 functions *f(t)* are selected according the value of *C*:

$$f(t) = a_s + (a - a_s)t \quad \text{(linear) (1)}$$

$$f(t) = a_s + \begin{cases} (a - a_s)2t & for\ t \le 0.5 \\ (a - a_s)2(1-t) & for\ 0.5 < t \le 1 \end{cases} \quad \text{(triangle) (2)}$$

$$f(t) = (2t^3 - 3t^2 + 1)a_s + (-2t^3 + 3t^2)a + (t^3 - 2t^2 + t)g_s \quad \text{(Hermite function) (3)}$$

with time *t*_ [0,1], the amplitude a_s at the beginning of the FAP at *t*=0, and the gradient g_s of *f*(0) which is the FAP amplitude over time at *t*=0. If the transition time *T*_1, the time axis of the functions (1) to (4) has to be scaled. These functions depend on a_s, g_s, *a* and *T*, and thus they are completely determined as soon as the FAP bookmark is known.

The Hermite function of third order enables one to match the tangent at the beginning of a segment with the tangent at the end of the previous segment, so that a smooth curve can be guarantied. Usually, the computation of the Hermite function requires 4 parameters as input, which are a_s, g_s, *a* and the gradient of *f*(*t*) at *t*=1. Here a horizontal gradient at the end of the transition time is assumed. The gradient $g_s(t)$ at time *t* is computed according to

$$g_s(t) = (6t^2 - 6t)(a_s - a) + (3t^2 - 4t + 1)g_s \quad (4)$$

with a_s, g_s , and a defined by the amplitude prior to the current bookmark.