# List from meeting

* Exclude marker
* ~~Initial position of model (calculated using markers)~~
* ~~Initial segment scaling (calculated using markers)~~
* ~~Range of motion for joints (to assist kinematics)~~
* ~~GRF prediction~~
* ~~Model independent of windows folder structure~~
* ~~Model independent of AnyScript folder structure (position of HumanModel and EnvironmentModel, C3DFileData etc)~~
* ~~tStart/tEnd perturbation to allow calc of Vel and Acc in the inverse dynamic model~~
* Parameters of #define ?????
* Marker drup-up handling?????
* ~~Marker renaming vs. standard protocols~~
* ~~Single Model-load-step~~
* ~~Separation of Trial/subject/lab specific data.~~
* ~~Fusion of AnyGait/Mocap~~
* Optimization of bodypart without marker ?????? (ST)
* ~~Multiple calibration trials ( standing, dynamic reference)~~
* ~~Study to just visualize the C3D data~~
* Comparing trials ??????
* ~~Easy to read and understand model ( hiding vs. showing the complexity of model)~~
* ~~Residual reduction algorithm to segment parameters (mass/inertia)~~
* ~~More environment directly created from the c3d files.~~

# ======================================================== Structured in groups

**New model features (in anyscript)**

* GRF prediction
* Study to visualize markers
* Range of motion for joints
* Initial segment scaling / position from markers
* Residual reduction algorithm for segment parameters.
* Use default arguments in ClassTemplates
* Overwrite lowpass filter for individual markers
* Create environments from C3D files

**New Featrures in AnyBody**

* Make it unnecessary to consider ApproxVelAccPerturb when setting tStart/tEnd
* Delay reading joint angles from files until it should be used.
* New way of transferring kinematics from the kinematics model to the inverse dynamic model
* Fix bug in

**Mocap model customization to specific usecases**

* Marker definition for many marker protocols.
* Fusion with AnyGait
* Multiple calibrations trials (standing and dynamic reference)

**General structure/appearance of the model**

* Make it easier to understand the model ( especially for beginners)
* Model independent of windows folder structure
* Model independent of HumanModel/Environment model
* Separate Trial/subject/ lab specific data
* Fusion with AnyGait

# Mocap model issues:

## Transferring joint angles:

Files: JointAngleOutputs.any, JointsAndDriversOptimized.any

Problems:

* Model does not load unless the files exist. This is a real problem with the combined model where both kinematics and inverse dynamics are in one model.
* Model has to be reloaded after joint angles are save to the file
* Adding new body parts or other things to the model is a pain.

## Model should work without specifying a special AnyMan.any file in the model model config

## BodyModel\_no\_muscles\_hack.any should handle if body parts are excluded.

## Making class templates independent of hard-coded model structure.

Files: CreateMarkerDriverClassNew.any, OptimizeAnthropometricsOnOff.any

Bug/feature in anybody can be used to accomplish this the following is a valid

nStep= Main.ModelSetup.nStep.Main.ModelSetup.nStep

The parser restarts if it meets an Main keyword

## Filter specification should work when no analog data is included in the model

Model should load even if no markers are included and no segments are optimized.  
ERROR(OBJ3) : C:/U..s/M..n/D..s/G..b/a..p/B..l/BaseModel\_Both.any(23) : ParameterIdentification : Member expectations are not satisfied :

Members of class AnyDesVar or derived are expected in AnyOptKinStudy : 0 found, 1..Inf expected.