

Computational Biomechanics

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General



- The AnyBody Modeling System is..
 - MusculoSkeletal Modeling System
 - analyzing reactions in the human body
 - between the human body and an environment.



- Environment can be
 - something within (implant, e.g. knee or hip device),
 - something attached to (exoskeleton, e.g. knee brace or spacesuit) or
 - something interacting (e.g. automotive seat, wheelchair, ...)

Inverse Dynamics of Muscle Systems





http://www.ebay.com/itm/Dynamometer-130kg-Measure-Hand-Grip-Gym-Measurement-Power-Strength-Dynamometre-/152408324194?hash=item237c3e7862;g:K30AAOxyP4dTcdFS

Anybody technology





- C f = r
 - -f muscle or joint forces
 - -r known forces (internal & external)
 - C matrix of equation coefficients

- Muscle forces $f \ge 0$
- Muscle redundancy



Linear Muscle Recruitment



$$G = \frac{f_1}{N_1} + \frac{f_2}{N_2}$$

• With *N* as normalization factor (muscle activity)

$$\begin{array}{ll} Min & G(f^M) \\ s.t. & C f = r \\ f^M \ge 0 \end{array}$$

Quadratic Muscle Recruitment



$$G = \sum_{i} \left(\frac{f_i}{N_i}\right)^2$$

$$\begin{array}{ll} Min & G(f^M) \\ s.t. & C f = r \\ f^M \ge 0 \end{array}$$

Polynomial Muscle Recruitment



$$G = \sum_{i} \left(\frac{f_{i}}{N_{i}}\right)^{p}$$

 $\begin{array}{ll} Min & G(f^M) \\ s.t. & C \ f = r \\ f^M \geq 0 \end{array}$





$$G = \max\left(\frac{f_{\rm i}}{N_{\rm i}}\right)$$

- \rightarrow linear problem
- \rightarrow nummerically efficient
- →physiologically reasonable



Mechanical model

- Remind: C f = r
- Segment *i*

$$\begin{aligned} q_i &= \begin{bmatrix} r_i \\ p_i \end{bmatrix} & \text{Position} \\ 4 \text{ Euler parameters} \\ \nu_i &= \begin{bmatrix} \dot{r_i} \\ \omega_i \end{bmatrix} & \text{Translation velocities} \\ \text{Rotation velocities} \end{aligned}$$

- $\rightarrow \phi(q,t) = 0$
- Kinematic equations \rightarrow obtain q, v, \dot{v}

Dynamic equilibrium

$$\begin{bmatrix}
m_i I & 0 \\
0 & J_i
\end{bmatrix} \dot{v}_i + \begin{bmatrix}
0 \\
\overleftarrow{\omega}_i' J_i' \omega_i'
\end{bmatrix} = f_i$$
• f:
- Muscle
- Reaction
$$\begin{bmatrix}
f \\
- Reaction
\end{bmatrix} f \\
- Cf = r \\
- Applied] r$$

Structure







Propeties: Mass, reference node...

Loads

• ANYSCRIPT



Main = {
 // The actual body model goes in this folder
 AnyFolder MyModel = {
 // Global Reference Frame
 AnyForeKefFrame GlobalRef = {
 AnyForeKefFrame drw = {
 ScaleMYZ = (1,1.1)/10;
 RGB = {(1,0.0};
);
 };
 AnyRefNode M1 = {
 sRel = {0, 0.05, 0};
 };
 AnyRefNode M2 = {
 sRel = {0, 0.1, 0};
 };
 // Global reference frame
 // Just a simple arm segment hinged at its left end
 AnySeg Seg = {
 Mass = 2;
 Jij = {1, 1}/1000;
 Just = {1, 1}/1

Picture Sources: https://anyscript.org/assets/images/teaser.jpg

AMMR



repository of musculoskeletal models

\rightarrow Fast use



Shoulder

118 muscle fascicles on each side
 Wrapping of muscles by contact mechanics
 Contact criterion in the GH joint
 Veger et al 1991. Biomech. 24, 615.29
 Van der Heim 1994 J. Biomech. 27, 551.59
 Veger et al 1997. Biomech. 30, 647.52



TS Scapula thoracic gliding plane, ellipsoid Al Scapula thoracic gliding plane, ellipsoid

AC Spherical joint GH Spherical joint SC Spherical joint











• Demo