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- Observations:
- Tubing and Quaternion Frame Space. Any path of frames on this space can be used to solve the *tubing problem*.
- Minimality. The PT frame appears to be unique frame with minimum total rotation.
- desired boundary twist uniformly across vertex frames: This is best done using uniform Quaternion distances between uniformly spa-Distributed Twist. A conventional compromise distributes a user-
- tially sampled frames.
- **Hybrids.** On *closed curves*, Frenet frame is periodic, PT is not. Add fixed angular increment throughout to make PT periodic.
- Initial angular velocity. Can give the frame an arbitrary number of twists using  $\sigma \neq 0.$  Minimal tangential acceleration version corresponds to quaternion treatment by Barr, Currin, Gabriel, and Hughes (Siggraph 92).



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# Example: Quaternion Protein Frame Statistics



of alternative geometries. (right) Quaternion maps showing the orientation tries for the protein YvyC from Bacillus subtilis, 2HC5. (left) The collection space geometry spreads for each individual amino acid. Quaternion maps for NMR data describing 10 different observed geome-

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### **Basic Background: Orientation Frames in** Bioinformatics

- Proteins are important. The entire machinery of life depends on metabolism.
- arranged in a sequence, but with very complicated 3D geometry. or even thousands, of amino acids Proteins are long chains of frames. Proteins consist of hundreds, on frames
- Traditional orientation tools describing proteins are primitive. The Ramachandran plot relates amino acid n to amino acid  $n \pm 1$ - that's it!
- Ramachandran statistics are impossible. With only local informanon-rigid protein orientation distribution. tion, you can't compare distant active sites, or gather statistics on

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# New Progress: Quaternion Frames in Proteomics

- orientation frame labels. data to construct precise, amino-acid-residue by amino-acid-residue, The PDB has massive protein geometry data. We can mine that
- PDB geometry to quaternion frame sequences. Amino acid quaternion frames. It is straightforward to convert the
- Using our quaternion display tricks, global information about residue alignment is directly visualizable.
- Our just-published JMGM paper applies quaternions to many
  proteomics problems. For additional information, see A. Hanson nion Maps of Global Protein Structure." (Fall 2012). and S. Thakur, Journal of Molecular Graphics and Modelling, "Quater-

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### **Basic Procedure**

- Library of 20 amino acids. Proteins link these together with peptide and joins together as C'–NH–C  $_{\alpha},$  kicking off a water, H  $_2O.$ bonds: a C'–OH unit on one end sees an  $\text{NH}_2\text{-}\text{C}_\alpha$  on the other side,
- Pick Three Atoms. Any three noncollinear atoms are sufficient to poses than others. define a quaternion frame, but some are more useful for specific pur-
- Compute Quaternion Frames for the whole protein.
- View frame sequence on the quaternion sphere. Global compar-isons as well as local comparisons can be made with a sequence of quaternion frames.
- Study the map. The map itself can be used to perform orientation statistics and similarities unobtainable by other methods



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native geometries. (b) Quaternion map clusters.



## Part II c: Dual Quaternions

- Quaternions Describe Only 3D Rotations. A computer tions and Translations. graphics scene must place elements using both Rota-
- Worse: Quaternion "Vector" Is Wrong. The "vector" quaternions does not result from any sensible translationtion, not a vector (1989 paper by Altmann). (0, x) in pure like transformation.  $(0, \mathbf{x})$  in  $R \cdot \mathbf{x} = q \star (0, \mathbf{x}) \star q \star$  is a 180-degree rota-

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### Dual Quaternions...

- Brief History. Dual quaternions (biquaternions) were first investitreatments can be found, e.g., from the German school of Blaschke (1960), and are used in theoretical mechanics (Bottema and Roth, dual quaternions and Clifford algebras. 1979; McCarthy, 1990), and in robotics. See also See Dorst et al. gated by Clifford (1873), and elaborated by Study (1891). Modern Geometric Algebra for Computer Science for the connection between
- Resources for Graphics Applications. Kavan et al. (TOG, 2008) an excellent summary, but misses a couple of fine points that we will to graphics for skinning problems, etc. (Appendix of Kavan et al. has have spurred the transfer of dual quaternion methods from robotics look at below.)

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rotation into a translation.

Usual:  $i^2 = -1$ :  $e^{i\theta}$ II ||  $\frac{1+i\theta-\frac{1}{2}\theta^2-\frac{1}{3!}i\theta^3+\cdots}{\cos\theta+i\sin\theta}$ 

 $e^{i\theta}e^{i\phi}$ Ш  $\cos(\theta + \phi) + i\sin(\theta + \phi)$ 

 $= 0: e^{\epsilon t}$ II

Dual:  $\epsilon^2$  $e^{\epsilon x}e^{\epsilon t} =$  $1 + \epsilon(x+t)$ .  $1 + \epsilon t + 0$ 

Like the  $\theta \rightarrow 0$  limit of rotation. So the dual algebra looks like *small*  $\theta$  or *large radius* rotation, which is effectively translation.

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mann's objection is surprising New look for 3D Vectors: The key to removing Alt- Now we can make a vector from nothing using quater-• This gives us a way to make a True Vector. Then multiply the x by ε. • Replace the zero in (0, x) by a one nion conjugation:  $(1, \epsilon \mathbf{x}) = (1, \epsilon \mathbf{x}/2) \star (1, \epsilon 0) \star (1, \epsilon \mathbf{x}/2)$ . ... toward Dual Quaternions...  $\mathbf{x} = (1,0) + \epsilon(0,\mathbf{x})$  $\equiv$  (1,  $\epsilon \mathbf{x}$ ).



- Dual translation  $\tau(d)$ : Our new tool (with half-vectors) needed plus sign in sandwiched translation. dual conjugation  $\overline{a + \epsilon b} = a - \epsilon b$ . Need both to give
- is the dual quaternion  $\tau(\mathbf{d}) = (1,0) + \epsilon \left(0,\frac{\mathbf{d}}{2}\right) \equiv (1,\epsilon\frac{\mathbf{d}}{2})$
- Translate x to x + d by conjugate multiplication:  $\tau(\mathbf{d}) \star (1, \epsilon \mathbf{x}) \star \overline{\tau(\mathbf{d})^*} = (1, \epsilon (\mathbf{x} + \mathbf{d}))$

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### **Dual Quaternions Fix Hamilton**

 Rotation done right! SAME answer, of course, but now  ${\bf x}$  is no longer confounded with 180° rotation:

$$(1, \epsilon R \cdot \mathbf{x}) = q \star (1, \epsilon \mathbf{x}) \star \overline{q^*}$$

• complete OpenGL-style transformation,  $\mathbf{x}' = T \cdot R \cdot \mathbf{x} =$ Full SE(3) frame now possible! We can perform a  $R \cdot \mathbf{x} + \mathbf{d}$ , as:

 However, this is only half the story.  $\tau(\mathbf{d}) \star q \star (1, \epsilon \mathbf{x}) \star \overline{(\tau(\mathbf{d}) \star q)^*} = (1, \epsilon(R \cdot \mathbf{x} + \mathbf{d}))$ 

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## **Moving Centers with Dual Quaternions**

to use that representation for the FULL frame? Problem: what happened to  $(\cos, n \sin)$ ? Don't we want

• First step:  $T(r) \cdot R \cdot T^{-1}(r)$ . What happens to the fixedpoint rule? θ Ð

$$\tau(\mathbf{r}) \star \left(\cos\frac{\theta}{2}, \mathbf{n}\sin\frac{\theta}{2}\right) \star \tau(-\mathbf{r})$$
$$= \left(\cos\frac{\theta}{2}, (\mathbf{n} + \epsilon \mathbf{r} \times \mathbf{n})\sin\frac{\theta}{2}\right)$$

 $\bullet$  So the fixed point  ${\bf r}$  appears as a dual rotation axis  $\mathbf{r} \times \mathbf{n}$ , automatically in the plane perpendicular to  $\mathbf{n}$ .

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# Applications: Blending and Interpolation

 Blending for Skinning: Dual quaternions permit an unusually smooth combination of weighted skin vertices associated to two or more skeletal elements in character animation. The most rigorous methods are essentially dual quaternion extensions of the spherical centerof-mass methods of Buss and Fillmore (TOC, 2001). Faster, but less accurate methods, use the concept of Phong shading, renormalizing a linear combination of data sets (Kavan et al., TOC 2008).

Interpolation: Blending is a static process, and needs to be done to combine character body elements such as skin vertices at each moment. Interpolation for simulating moving object kinematics and controlling camera motion can also be accomplished by extending standard quaternion interpolation techniques to dual quaternions, though challenging issues such as how to control dual parameters and how to match rotational and translational speeds in a single interpolation introduce additional complexity and possible artifacts.

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## FINAL TUTORAL SUMMARY

- Quaternions nicely represent frame sequences.
   TUBES: Curve frames ⇒ quaternion curves. Exploit
- PROTEINS: Amino acid residue coordinates 

   quaternion frame maps. Apply to global comparisons and statistical distributions.
- DUAL QUATERNIONS: (From Clifford, 1873.) Extend quaternion rotation algebra to include translations. Applications include blending for skinning in figure animation, robot arm motion planning, etc.

ftp://ftp.cs.indiana.edu/pub/hanson/Siggraph12QuatCourse/ 55

