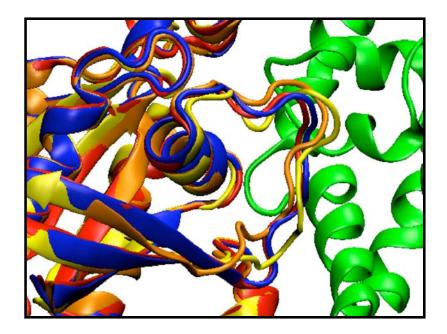
Protein Interface Modeling and Design with Kinematic Closure

Dan Mandell

RosettaCon August 5, 2009

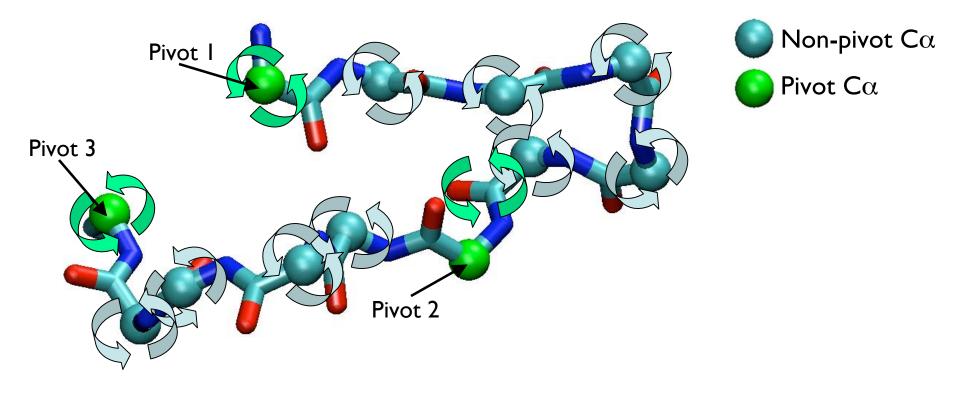
Conformational plasticity in protein recognition

Flexible interface loops facilitate binding of multiple partners, as in the Rac GTPase

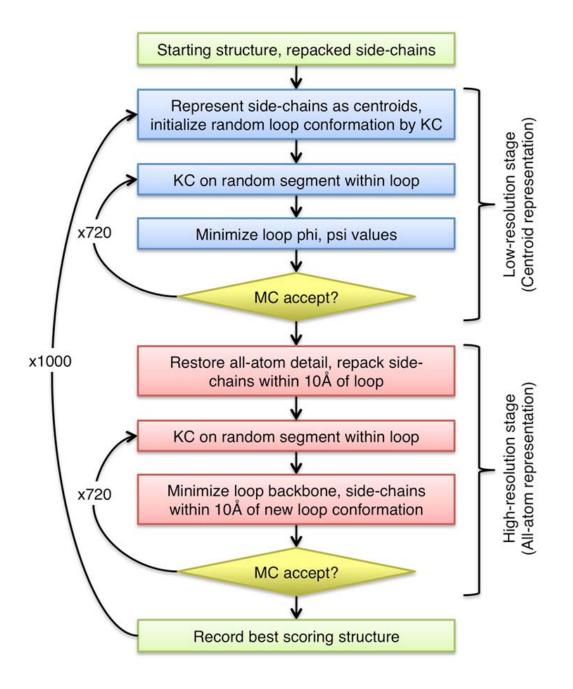


If we can accurately remodel protein interfaces, we can better engineer new functional protein interactions

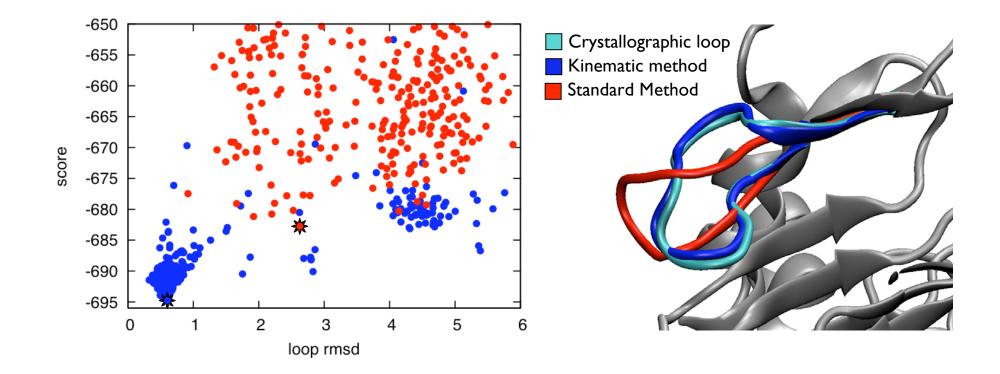
The kinematic closure (KC) move for peptide chains



Non-pivot torsions, sampled from Ramachandran space Pivot torsions, solved by closure Kinematic closure loop modeling protocol

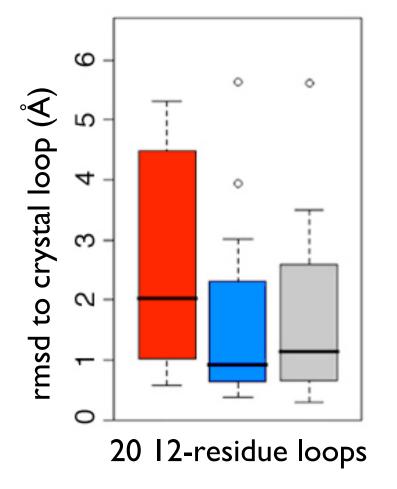


Near-native sampling by KC can yield high accuracy predictions



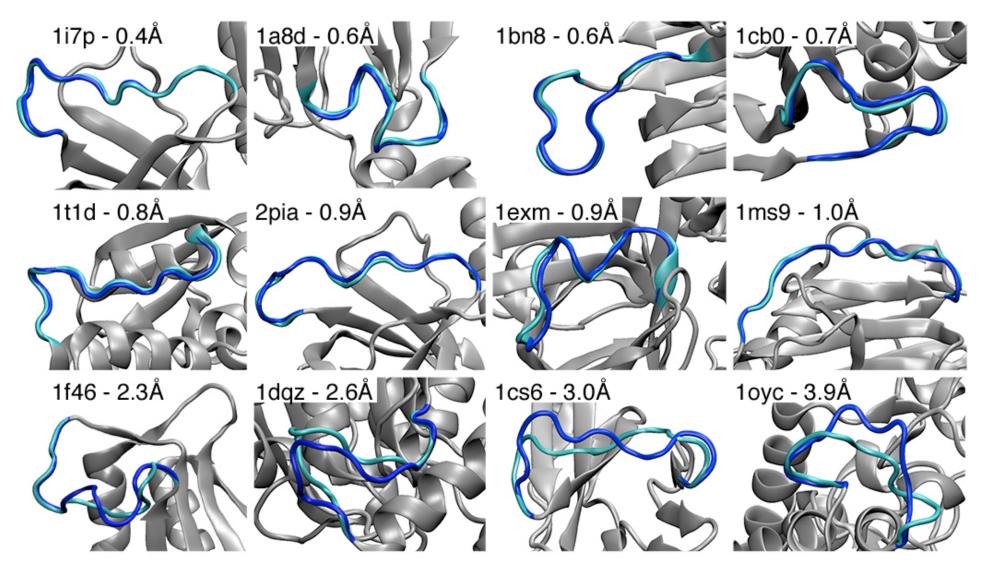
Performance across methods

Standard KC Molecular Mechanics



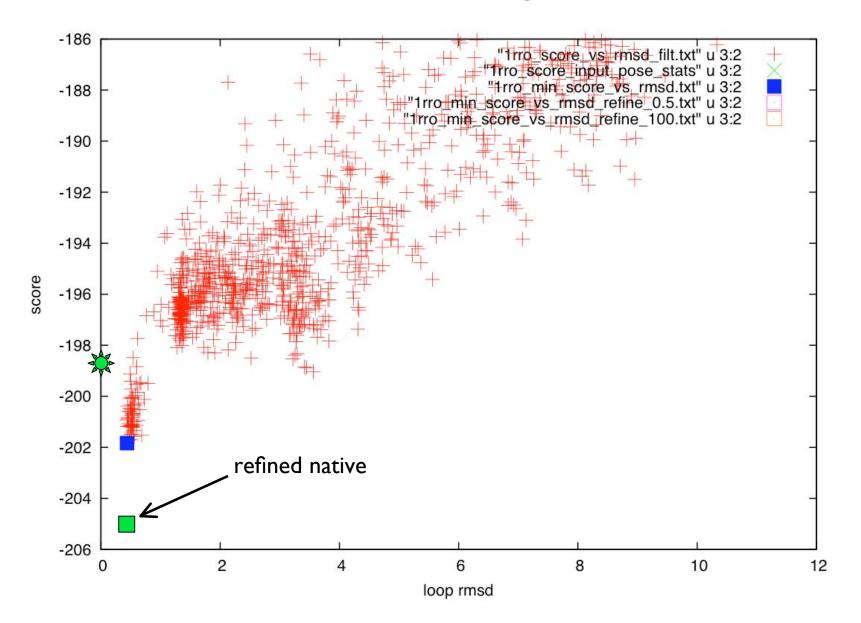
Mandell, Coutsias & Kortemme. (2009). Nat Methods 6(8):551-2

Representative set of reconstructions

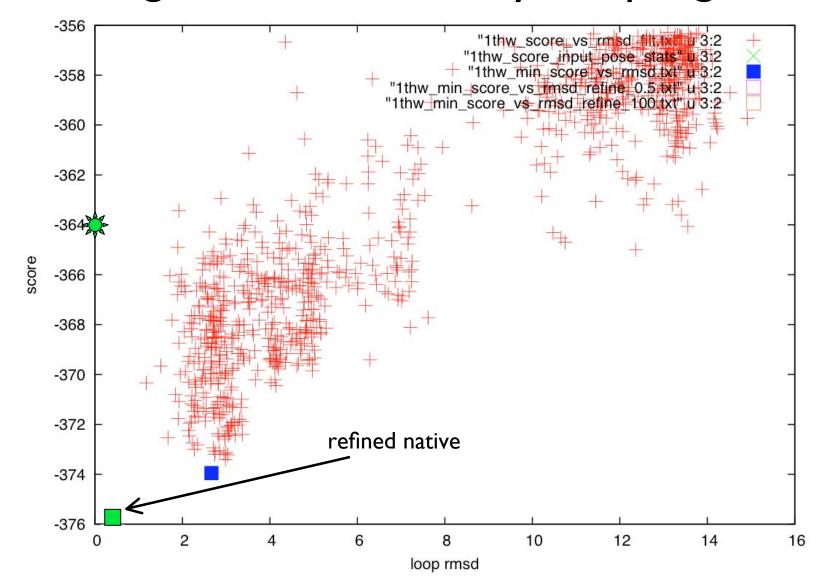


Mandell, Coutsias & Kortemme. (2009). Nat Methods 6(8):551-2

Effect of refining natives



KC can reveal that some purported scoring errors are actually sampling errors



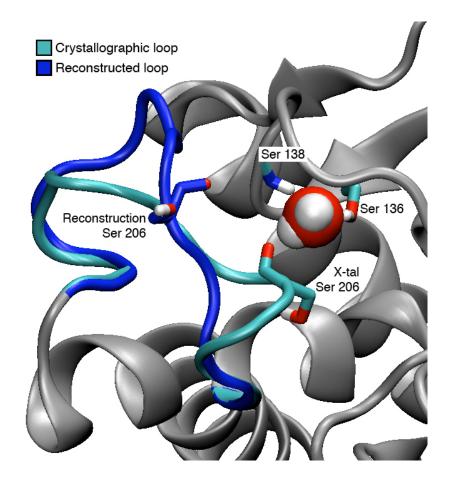
KC improves model accuracy by improved conformational sampling

KC Protocol		Standard Protocol	
Pdb	best scoring model score – crystallographic loop score	Pdb	best scoring model score – crystallographic loop score
1a8d	9.36	1a8d	21.98
1f46	7.91	1cnv	18.88
lcnv	5.25	1qlw	16.86
li7p	4.69	1dqz	16.66
1qlw	4.18	1bhe	14.89
1cs6	2.41	1f46	9.63
1my7	-0.56	1arb	6.32
2pia	-1.63	1bn8	6.12
1m3s	-3.89	1cs6	5.36
loyc	-5.58	1cb0	5.33
		1m3s	2.82
KC		1ms9	1.39
10 Failures		loyc	1.13
		1my7	0.61
6 Sampling		2pia	0.61
4 Scoring			

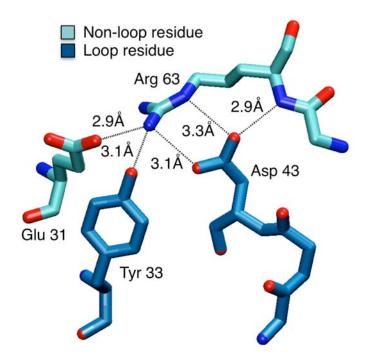
Standard 15 Failures

15 Sampling0 Scoring

Explicit water molecules not modeled

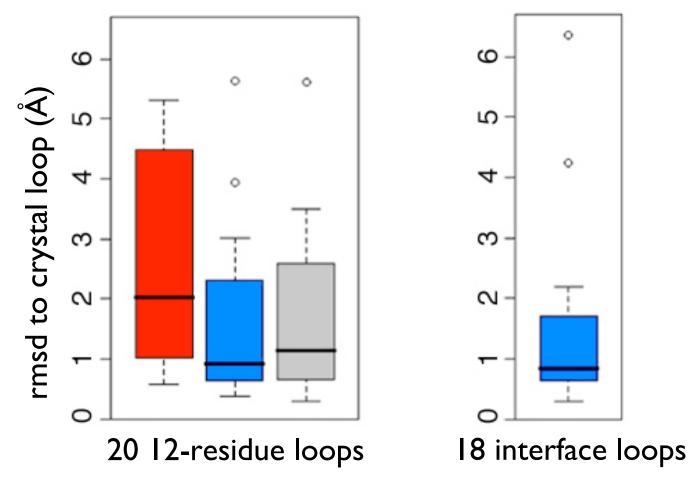


Complex hydrogen bonding networks can be difficult



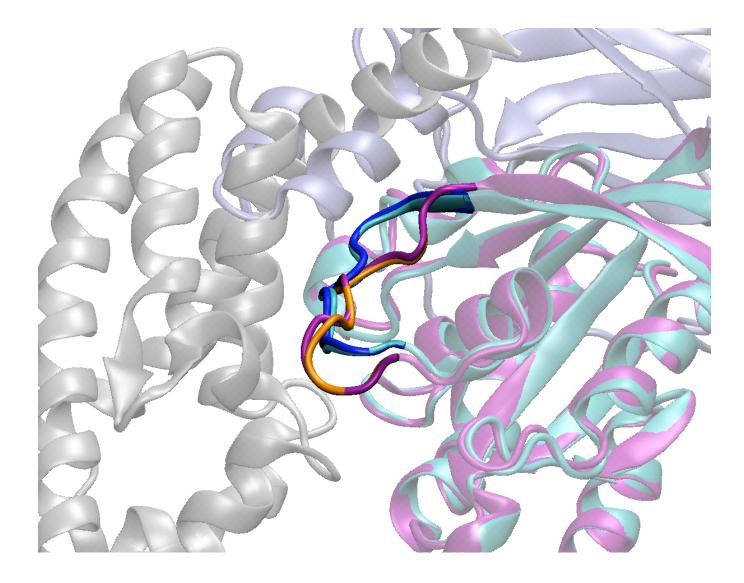
Conformational plasticity in interfaces

Standard 🔲 KC 🔲 Molecular Mechanics

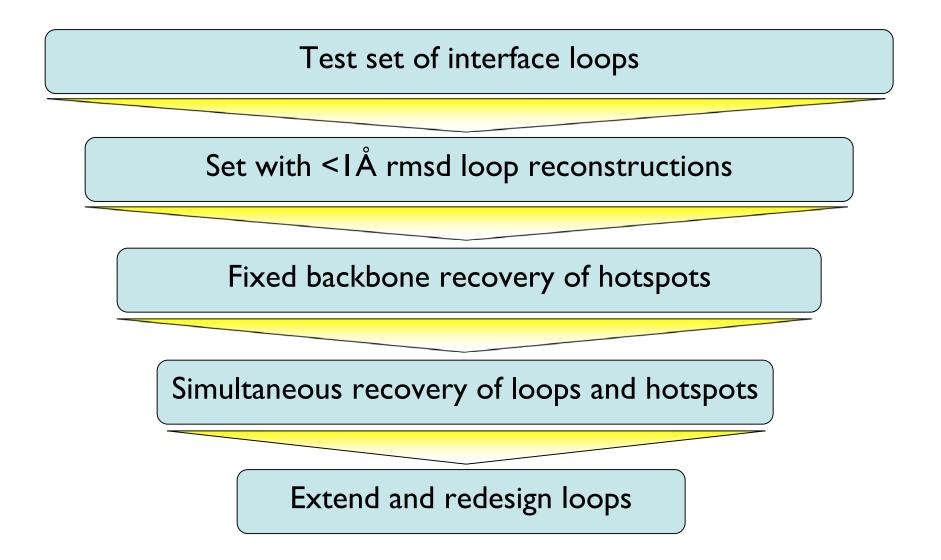


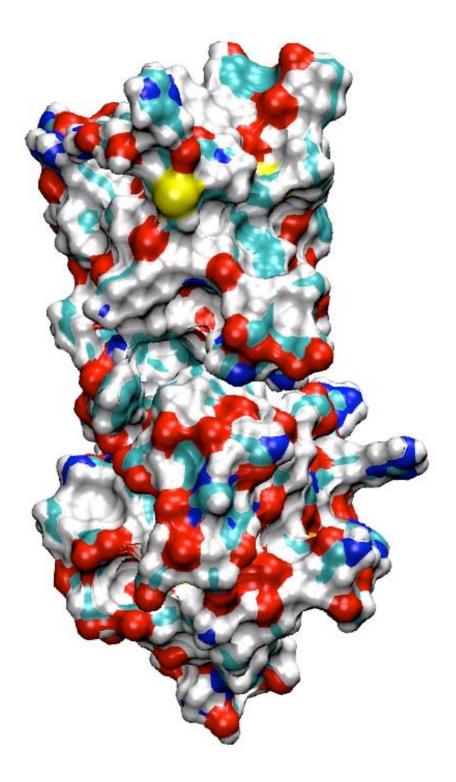
Mandell, Coutsias & Kortemme. (2009). Nat Methods 6(8):551-2

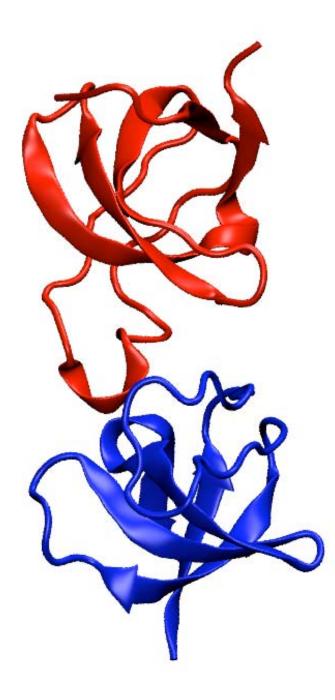
Conformational plasticity in interfaces

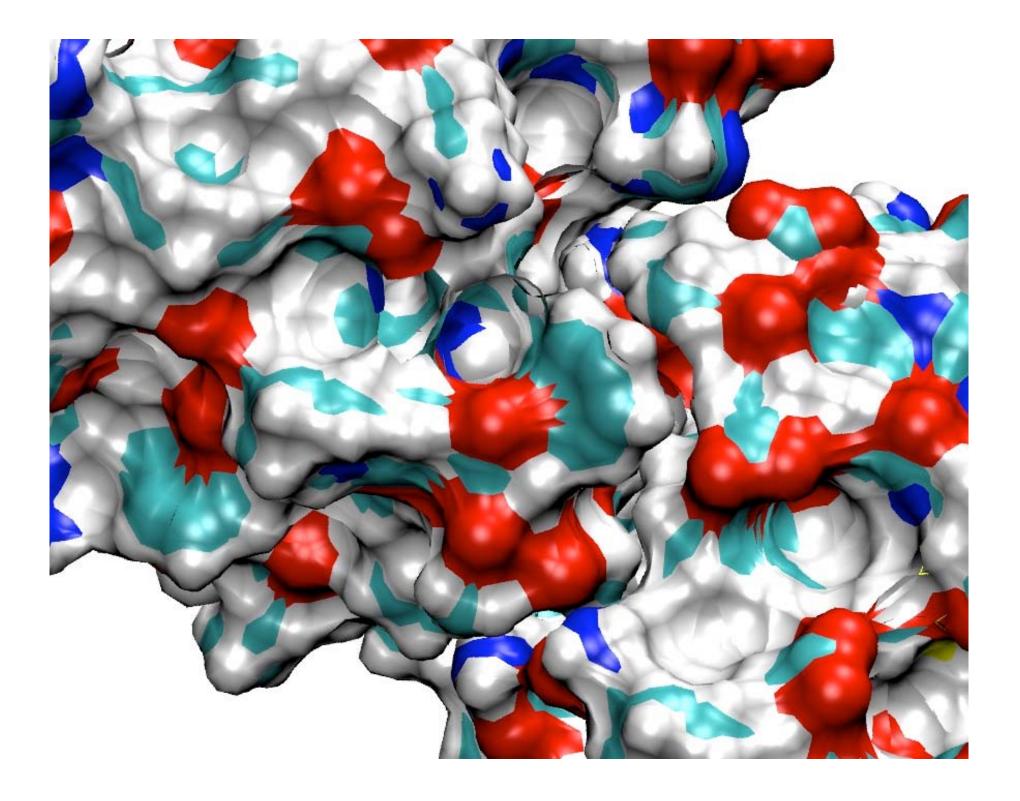


Kinematic modeling and design protocol









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