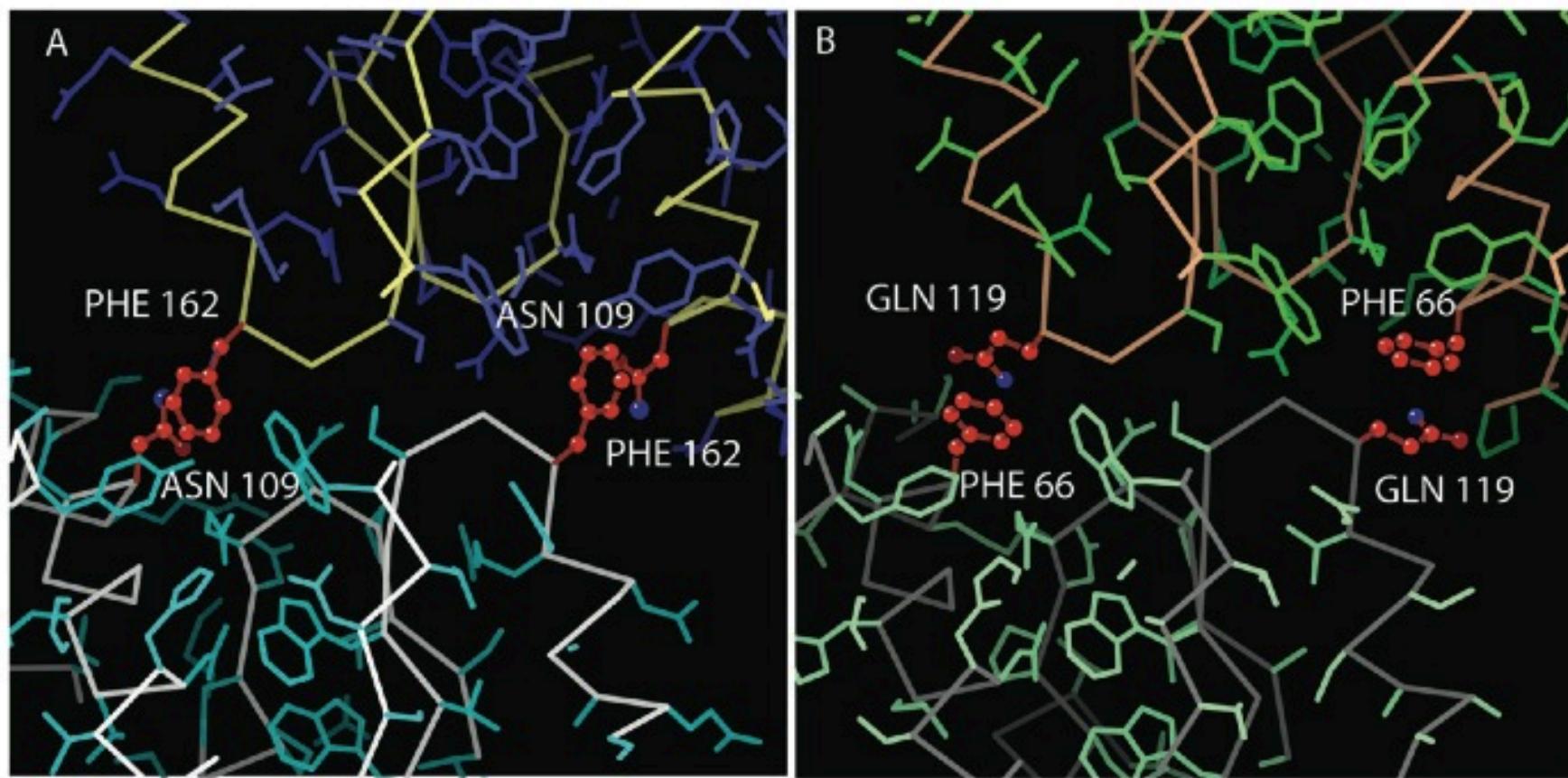


Coevolution between Proteins

- Protein–protein interactions effectively mediate molecular function and are maintained by the action of evolutionary pressure on the regions of the interacting proteins that contribute to binding.
- For the most part, selection restricts amino acid replacements, accounting for the conservation of binding interfaces. However, in some cases, change in one protein will be mitigated by compensatory change in its binding partner, maintaining function in the face of evolutionary change.



C

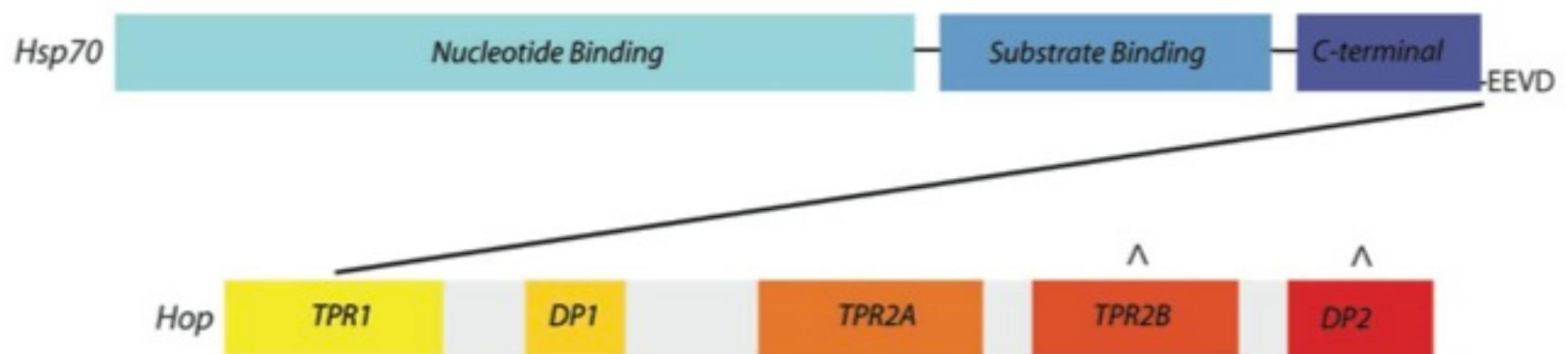
SODM_THET8	PQDIQTAVRNNGGGHLNHSLFWRLLT
SODF_NICPL	KGAPLPAF--NAAQAWNHQFFWESMK
SODF_SYNE7	GDASKAGLFNAAQAWNHSFYWNSIK
SODM_BACCA	PESIRTAVRNNGGGHANHSLFWTILS
SODF_METJ	PENLRGPVRNNGGGHANHSLFWKVLT
SODM_LISIV	PEEIRGAVRNHGGGHANHTLFWSSLS
SODM_MYCLE	--LNEKNLAFHLGGHVNSIWWKNLS
SODM_PROFR	--KLEKDLAFNLAGHVNVFWKNMA
SODF_METTH	--AALKELSFHVGGYVLHLFFWGNMG
SODM1_CAEEL	--ALQPALKFNGGGHINHSIFWTNLA
SODM_HUMAN	--ALQPALKFNGGGHINHSIFWTNLS

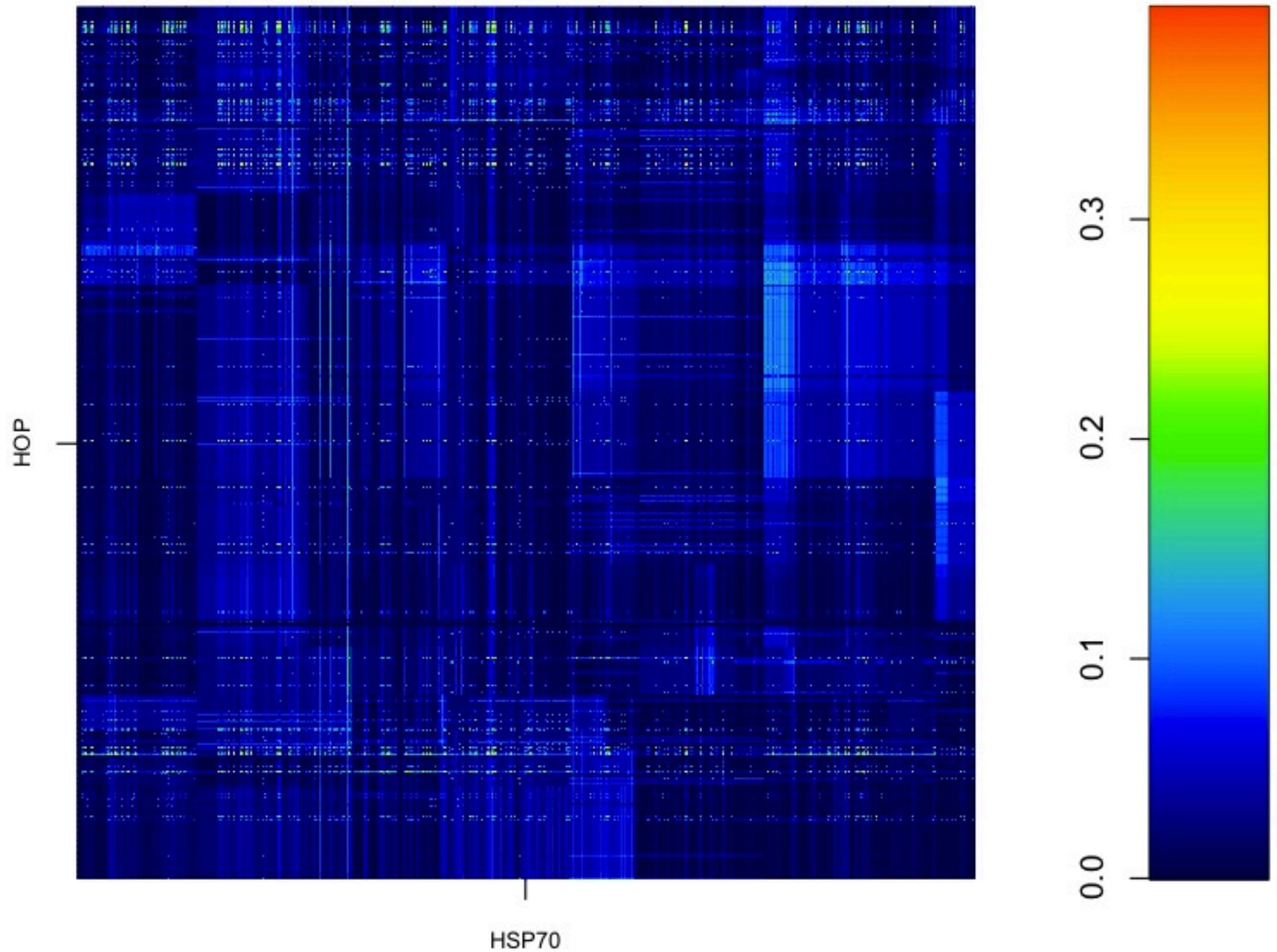
D

IDEQFGGGFQALKEKLTQAAAMGRFGSGWAHLVKD
INRDFGSYDAFVKEFKAAAATQFGSGWAHLAYK
IAADFGSFENFVTEFKQAAAATQFGSGWAHLVLD
INKKFGSFTAFKDEFSKAAAGRFGSGWAHLVVN
IKSDIGGLDTFKEAFTKAALTRFGSGWAHLSVT
IESEFGTFDEFKEKFNAAAAARFGSGWAHLVVN
IDETFGSFDKFRAQFSAAANGLOGSGWAHLGYD
IDEFFGSFDNMKAQFTAAATGIQGSGWAHLVWD
IEKDFGSFERFRKEFSQAAISAEGSGWAHLTYC
IKSDFGSLDNLQKQLSASTVAVQGSGWGWLGYC
IKRDFGSFDKFKEKLTAASVGVOGSGWGWLGFN

44 species ortholog alignments.

Example: Hsp70 & Hop





VVVVVV--VVV-VVV-VVVVV-VVVVVV-V-VVVVVVIIIIIII
DDDDDDDDDDDDDDDDDDDDDDDDDDDD-D-D-D---D-EEEEEEE

AAAAAA--AAA-AAA--AAAAA-AAAAAAA-AAAAAAGAASNNNNN
QQQQQQQQKQQQQQQQQQQQQQQ-Q-Q-Q-K-QTEDEEEE

AAAAAA--AAA-AAA--AAAAA-AAAAAAA-AAAAAAGAASNNNNN
EE-EEEEEEEEEEEEEEEEEEEEEEE-E-E-T-AEQQQQQ

AAAAAA--AAA-AAA--AAAAA-AAAAAAA-AAAAAAGAASNNNNN
KK-KKKKKKKKKTKKKKKKKKKKKKKKK-K-K-Q-KRRRRRR

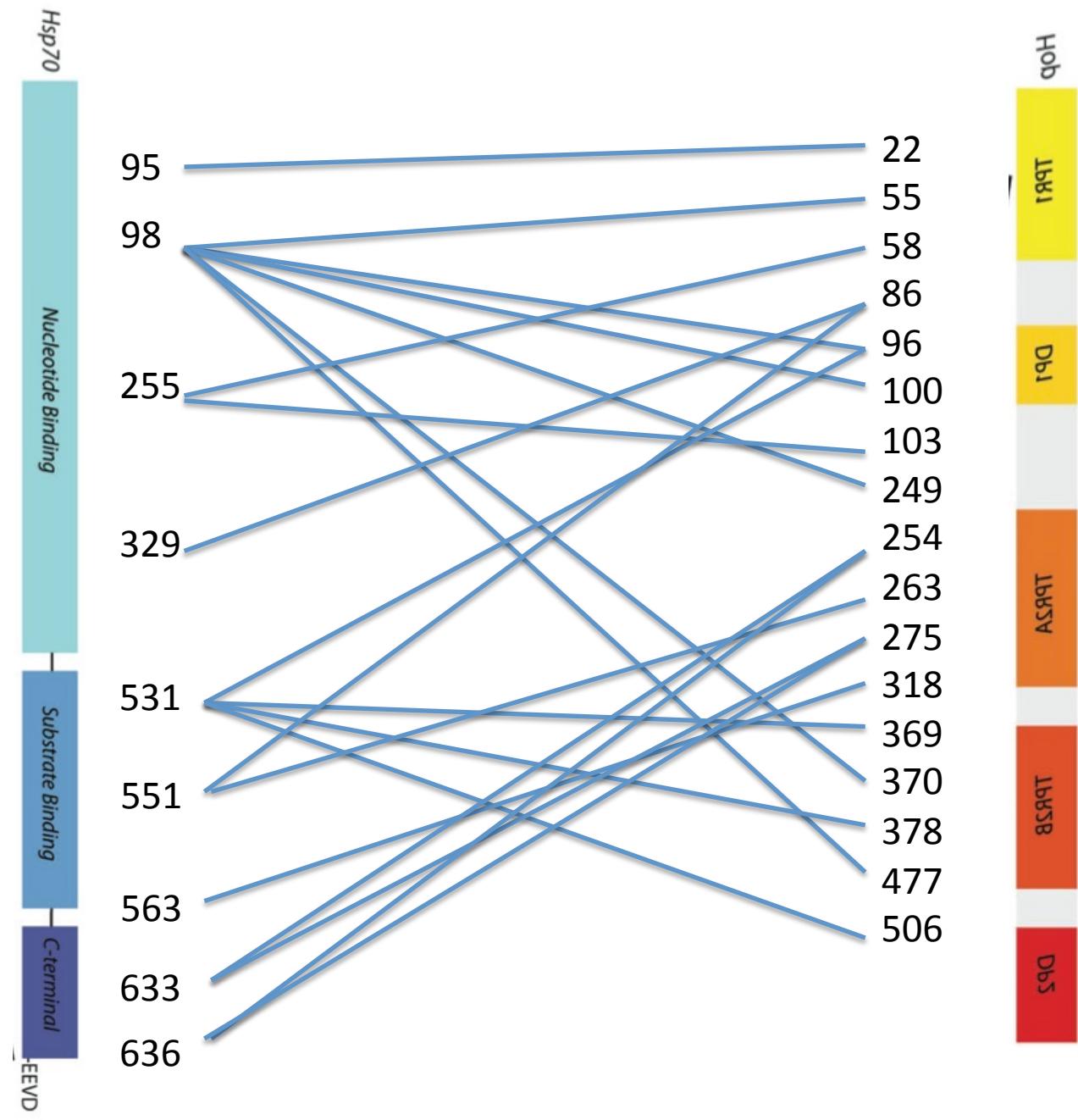
AAAAAA--AAA-AAA--AAAAA-AAAAAAA-AAAAAAGAASNNNNN
DDDDDDDDDDDDDDDDDDDDDDDDDD-DDDD-D-D-DGEEEEE

AAAAAA--AAA-AAA--AAAAA-AAAAAAA-AAAAAAGAASNNNNN
CCCCCC-CCC-CCCCCCCCCCCCCC-CCCCC-C-S-FSAAAAA

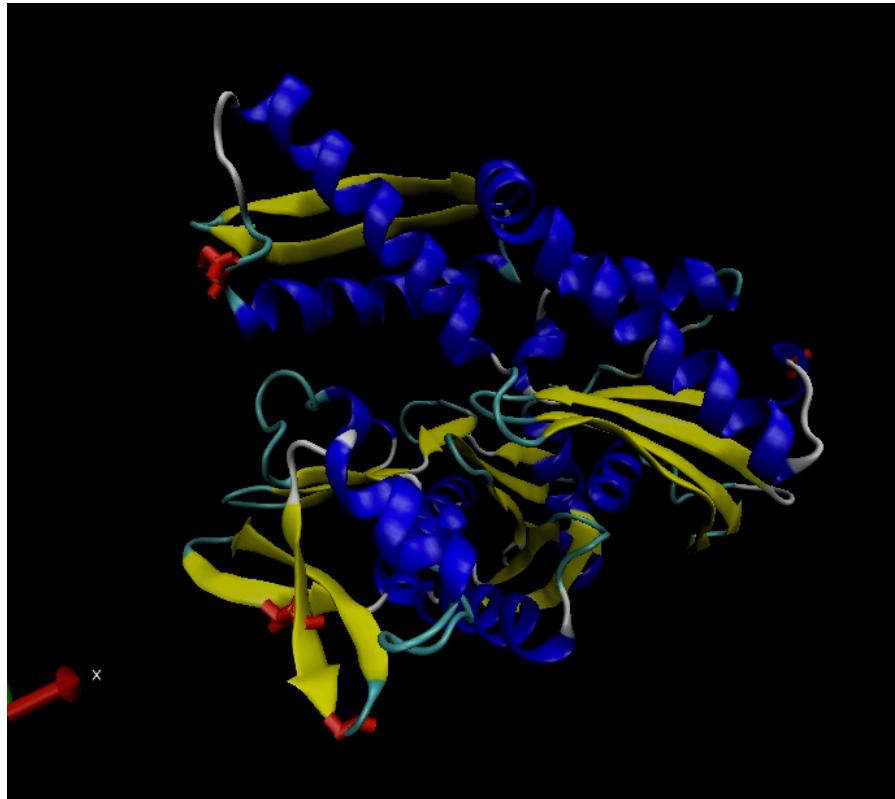
AAAAAA--AAA-AAA--AAAAA-AAAAAAA-AAAAAAGAASNNNNN
NNNNNN-NNNNNNNNNNNNNNNNNNN-NN-NN-N-T-NHTMTTT

EEEEEE-EEEEEEE-E---EEE-E-EE-EEEEEEEDDDDDDD
YYYYYYYYYFYYYYYYYYYYYYYY-Y-Y-Y-Y-L-LLLLLL

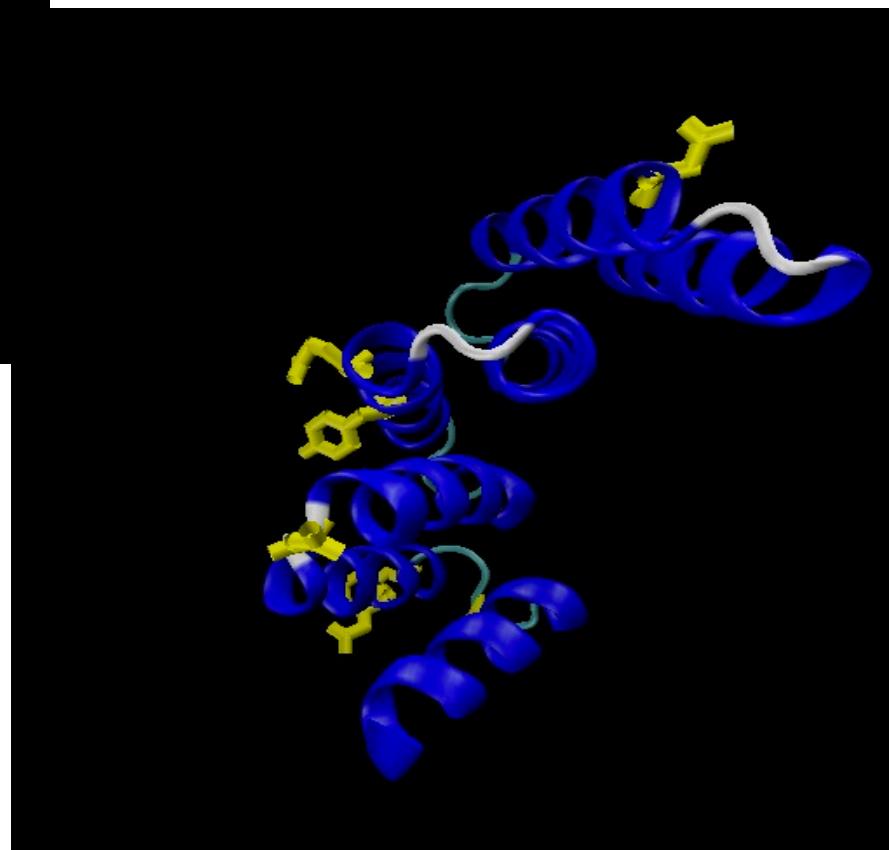
EEEEEE-EEEEEEE-E---EEE-E-EE-EEEEEEEDDDDDDD
AA-AAAAAAAAAAAAAAAA-A-A-S-PPPPPPP







HSP70



- The coevolutionary signal is subtle and somewhat weak in those proteins that only make transient interactions.
- Map 1000 genome SNP data & disease SNP data onto these pairs.