



Ubc12蛋白结构初步分析

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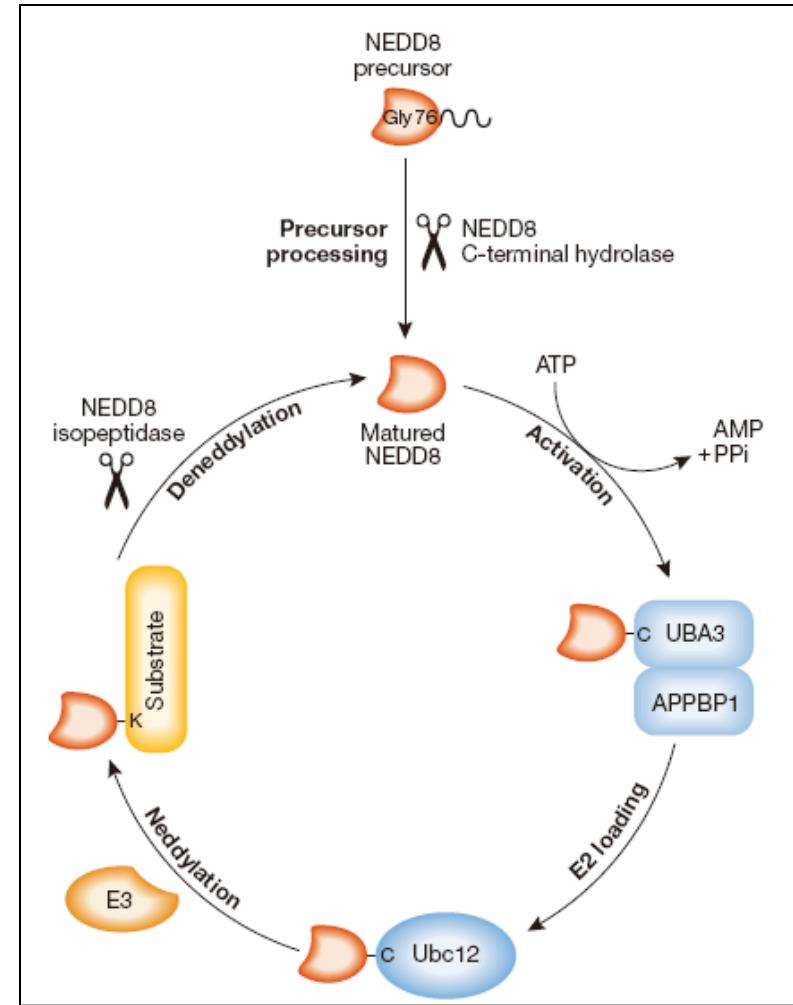
刁举鹏

Outline

- 1. Background on Neddylation and Ubc12
- 2. Goals of our project
- 3. Analysis by Bioinformatics tools
- 4. Conclusion

Background:

- NEDD8是一种类泛素化小分子，它在体内固有酶簇作用下被共价结合到底物蛋白上，参与蛋白质翻译后修饰，这一过程被称为Neddylation。



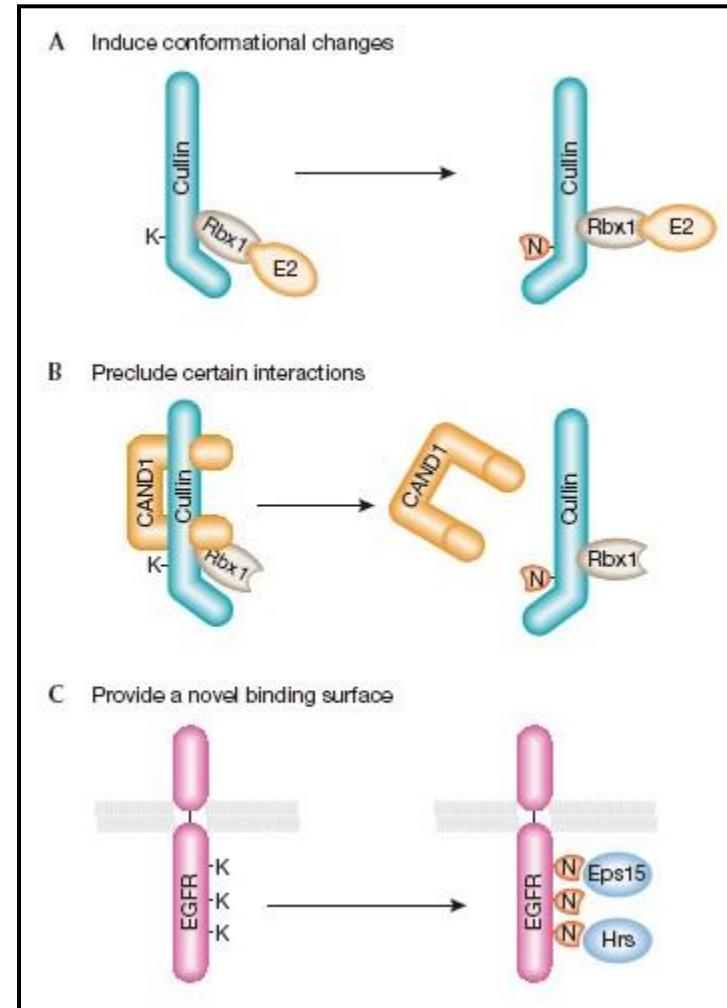
Neddylation过程 (Rabut G.et.al. 2008)

Background:

- Neddylation可以参与到细胞发育、细胞周期、信号转导等许多细胞调控。
- 目前发现的Neddylation的底物蛋白有：Cullin家族、p53、Mdm2、p73、表皮生长因子受体EGFR、核糖体蛋白家族等。
- E1：APPBP1–UBA3复合物
- E2：Ubc12（UBE2M）、UBE2F
- E3：Mdm2、Rbx1、Dcn1、SCF、c-Cbl等

Background

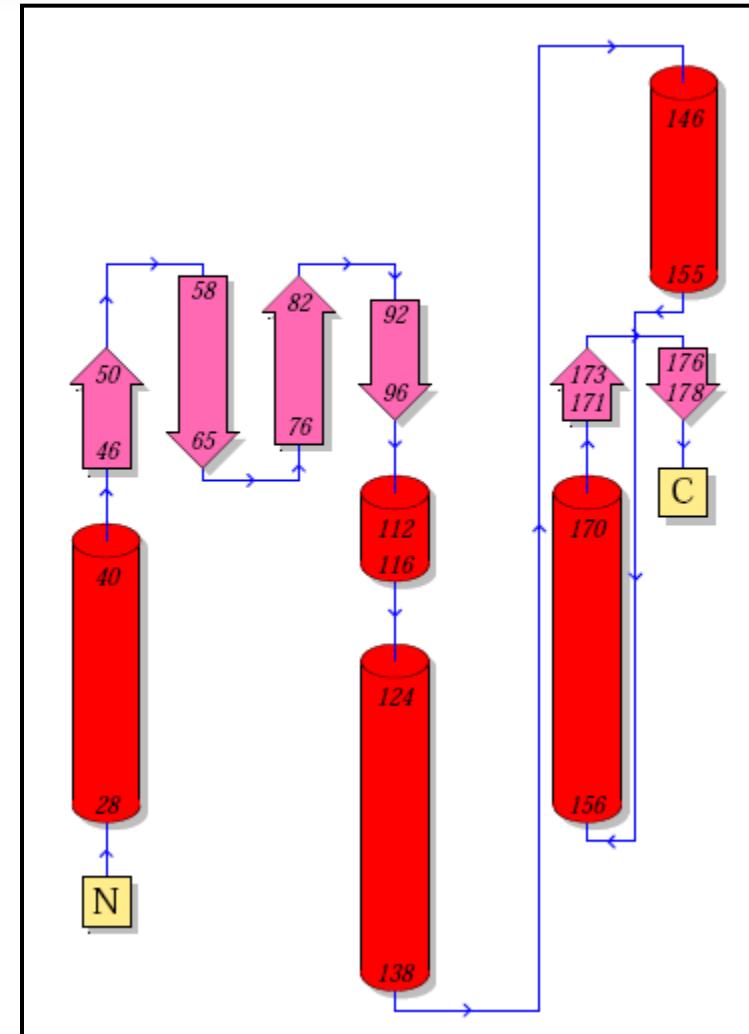
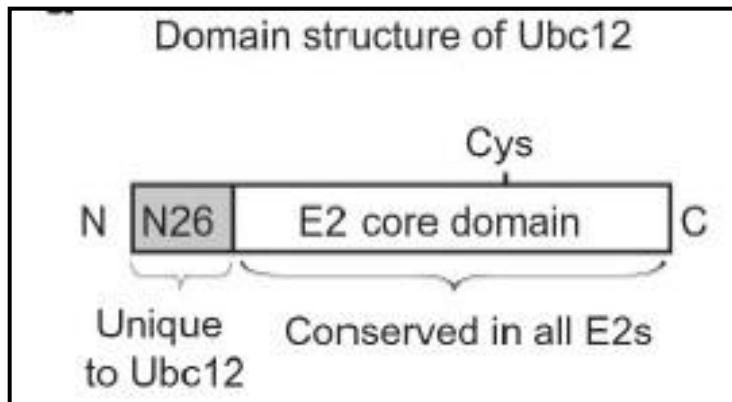
- 改变底物蛋白的构象
- 和其它相互作用蛋白竞争性结合
- 招募NEED8相互作用的蛋白。



Direct effects of neddylation (Rabut G.et.al. 2008)



- Ubc12由183个氨基酸残基组成，具有5个helix和6个strand。



Ubc12 Second Structure (<http://www.ebi.ac.uk>)

Goals of our project

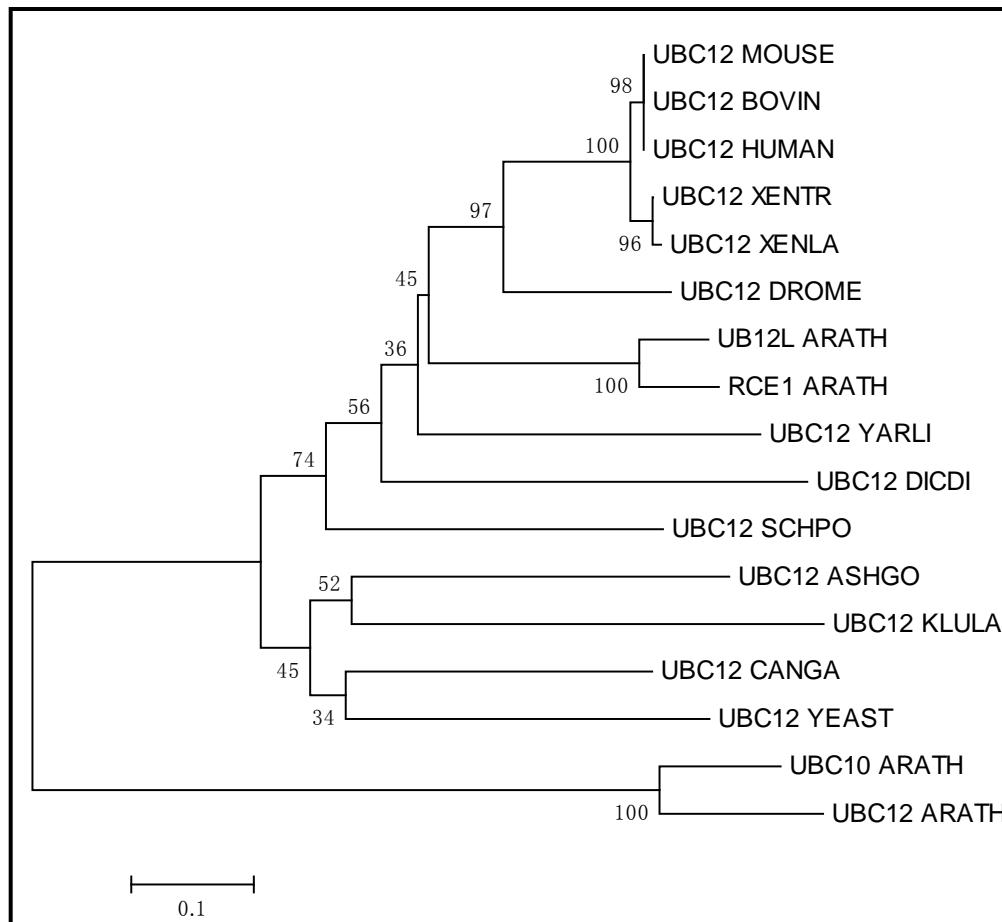
- 1. How does Ubc12 interact with E1?
- 2. How does Nedd8 transfer from E1 to E2?
- 3. How does Nedd8 transfer from E2 to substrate?



• 1.Sequence analysis

1	MIKLFSIKQQKKEEEAGGPKGSSKKASAAQLRIQKDINELNLPLKTCEIEFSDHDDLLNF	60	Q6P8D9	UBC12_XENTR
1	MIKLFSIKQQKKEEEAGGPKGSSKKASAAQLRIQKDITELNLPLKTCEIEFSDHDDLLNF	60	Q6DCZ9	UBC12_XENLA
1	MIKLFSIKQQKKEEEAGGPKGSSKKASAAQLRIQKDINELNLPLKTCDISESDPDDLLNF	60	A3KN22	UBC12_BOVIN
1	MIKLFSIKQQKKEEEAGGPKGSSKKASAAQLRIQKDINELNLPLKTCDISESDPDDLLNF	60	P61082	UBC12_MOUSE
1	MIKLFSIKQQKKEEEAGGPKGSSKKASAAQLRIQKDINELNLPLKTCDISESDPDDLLNF	60	P61081	UBC12_HUMAN
1	MIKLFTLKKQQKKDGEQKGSQQ---KKASAAQLRIQKDINELNLPLNTCATDFDPNDLLNF	57	Q9VSF3	UBC12_DROME
*****:*****: * . * . : *****:*****:*****:*****:*****:*****:*****:				
61	KLVICPDEGFYKGKGKFVFSFKVGQGYPHDPPKVKCETMVYHPNIDLEGNVCLNILREDWK	120	Q6P8D9	UBC12_XENTR
61	KLVICPDEGFYKGKGKFVFSFKVGQGYPHDPPKVKCETMVYHPNIDLEGNVCLNILREDWK	120	Q6DCZ9	UBC12_XENLA
61	KLVICPDEGFYKSGKFKVFSFKVGQGYPHDPPKVKCETMVYHPNIDLEGNVCLNILREDWK	120	A3KN22	UBC12_BOVIN
61	KLVICPDEGFYKSGKFKVFSFKVGQGYPHDPPKVKCETMVYHPNIDLEGNVCLNILREDWK	120	P61082	UBC12_MOUSE
61	KLVICPDEGFYKSGKFKVFSFKVGQGYPHDPPKVKCETMVYHPNIDLEGNVCLNILREDWK	120	P61081	UBC12_HUMAN
58	KLIISPDEGFYRDGRFVNFRVGSNYPHEPKVKCATQVYHPNIDLDGNVCLNILREDWN	117	Q9VSF3	UBC12_DROME
:.*****:.*:*****:*****:*****:*****:*****:*****:*****:*****:				
121	PVLTINSIIYGLQYLFLEPNPEDPLNKEAAEVLQNNRRLFEQNVQRSMRGYYIGSTYFER	180	Q6P8D9	UBC12_XENTR
121	PVLTINSIIYGLQYLFLEPNPEDPLNKEAAEVLQNNRRLFEQNVQRSMRGYYIGSTYFER	180	Q6DCZ9	UBC12_XENLA
121	PVLTINSIIYGLQYLFLEPNPEDPLNKEAAEVLQNNRRLFEQNVQRSMRGYYIGSTYFER	180	A3KN22	UBC12_BOVIN
121	PVLTINSIIYGLQYLFLEPNPEDPLNKEAAEVLQNNRRLFEQNVQRSMRGYYIGSTYFER	180	P61082	UBC12_MOUSE
121	PVLTINSIIYGLQYLFLEPNPEDPLNKEAAEVLQNNRRLFEQNVQRSMRGYYIGSTYFER	180	P61081	UBC12_HUMAN
118	PVLNINSIVYGLQFLFLEPNPEDPLNKEAADVLQTNRRQFENNWKAMRGGCCVGETYFEC	177	Q9VSF3	UBC12_DROME
*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:*****:				
181	CLK- 183 Q6P8D9 UBC12_XENTR			
181	CLK- 183 Q6DCZ9 UBC12_XENLA			
181	CLK- 183 A3KN22 UBC12_BOVIN			
181	CLK- 183 P61082 UBC12_MOUSE			
181	CLK- 183 P61081 UBC12_HUMAN			
178	CLK 181 Q9VSF3 UBC12_DROME			
**				

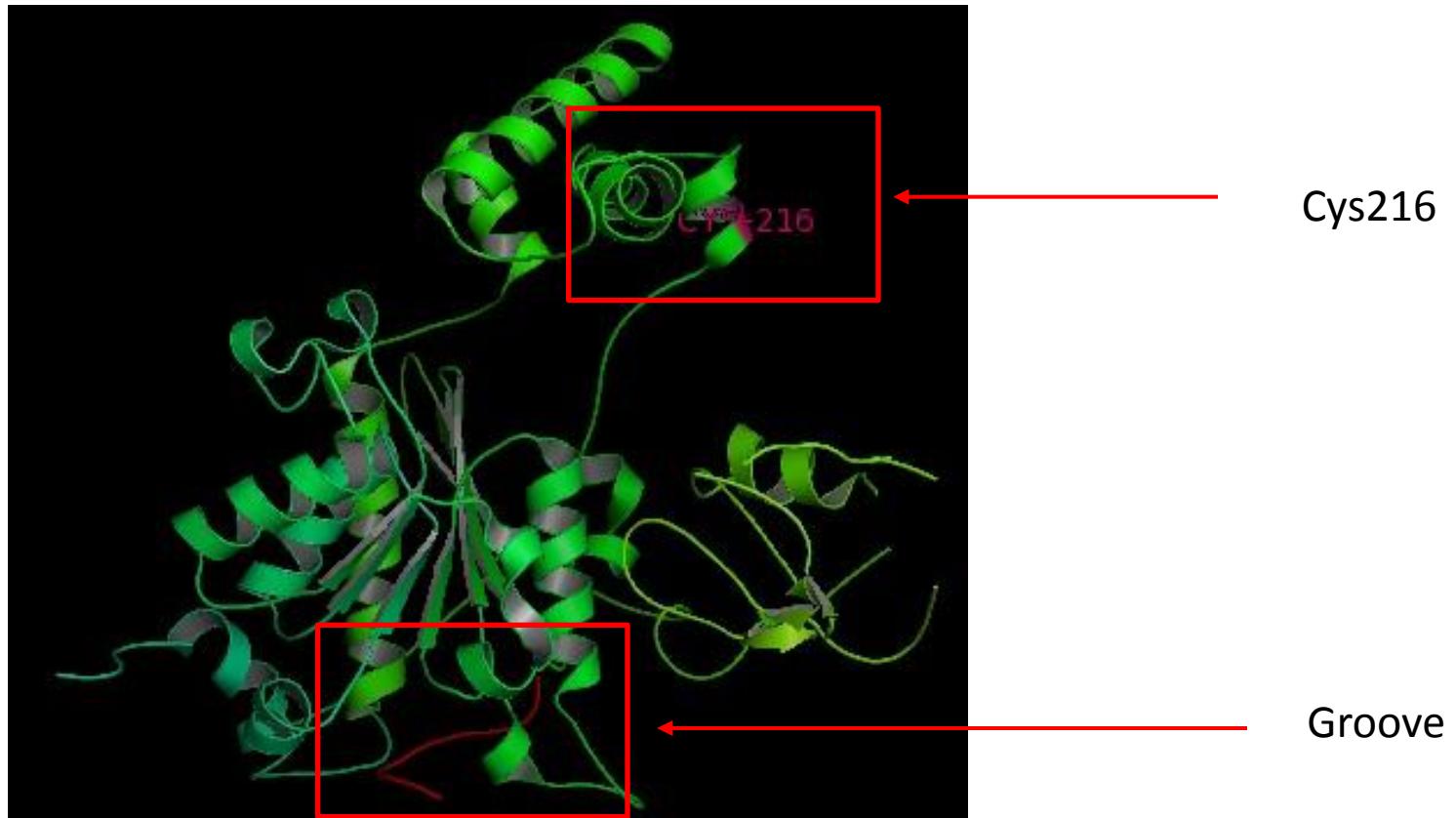
- 1.Sequence analysis



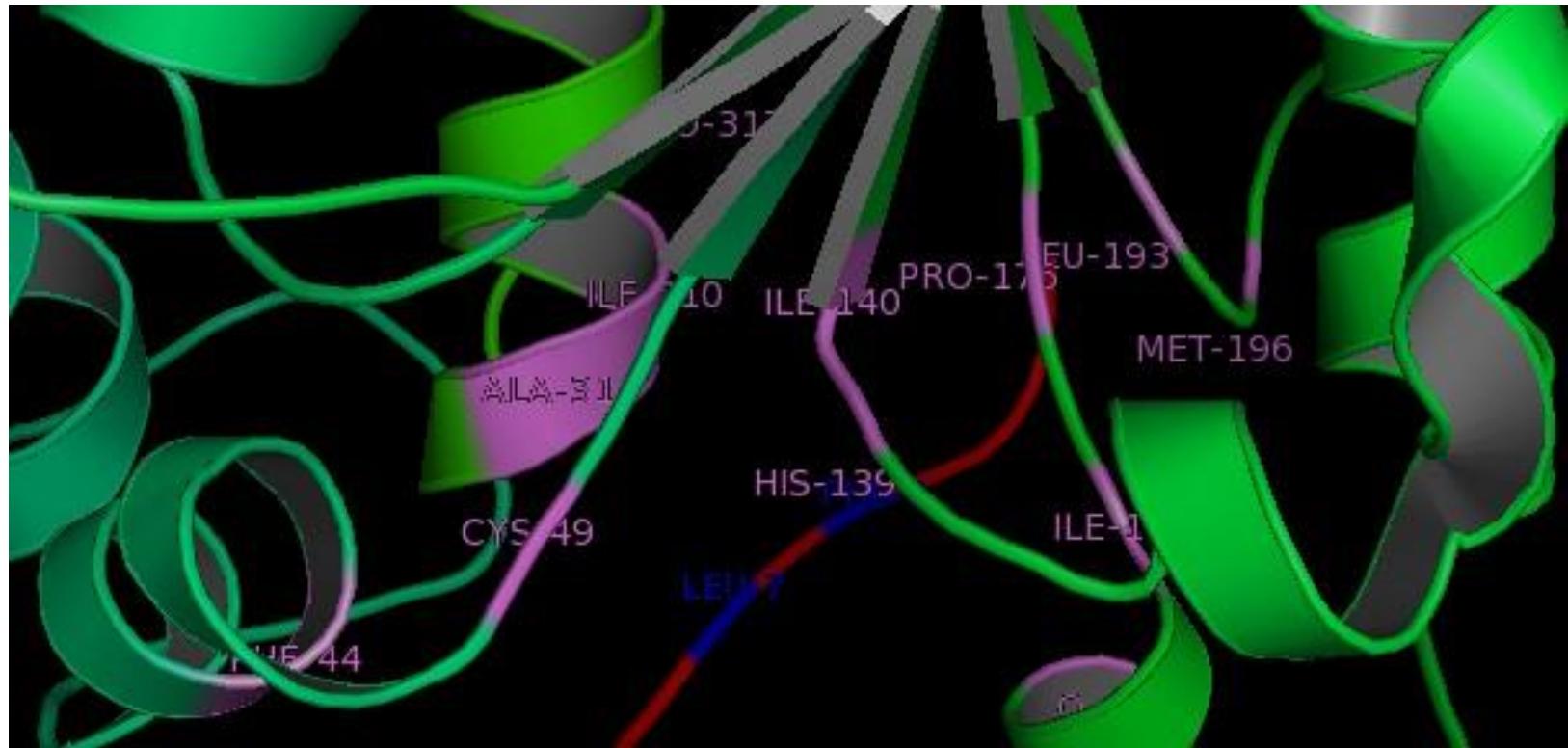
Ubc12直系同源系统发育树



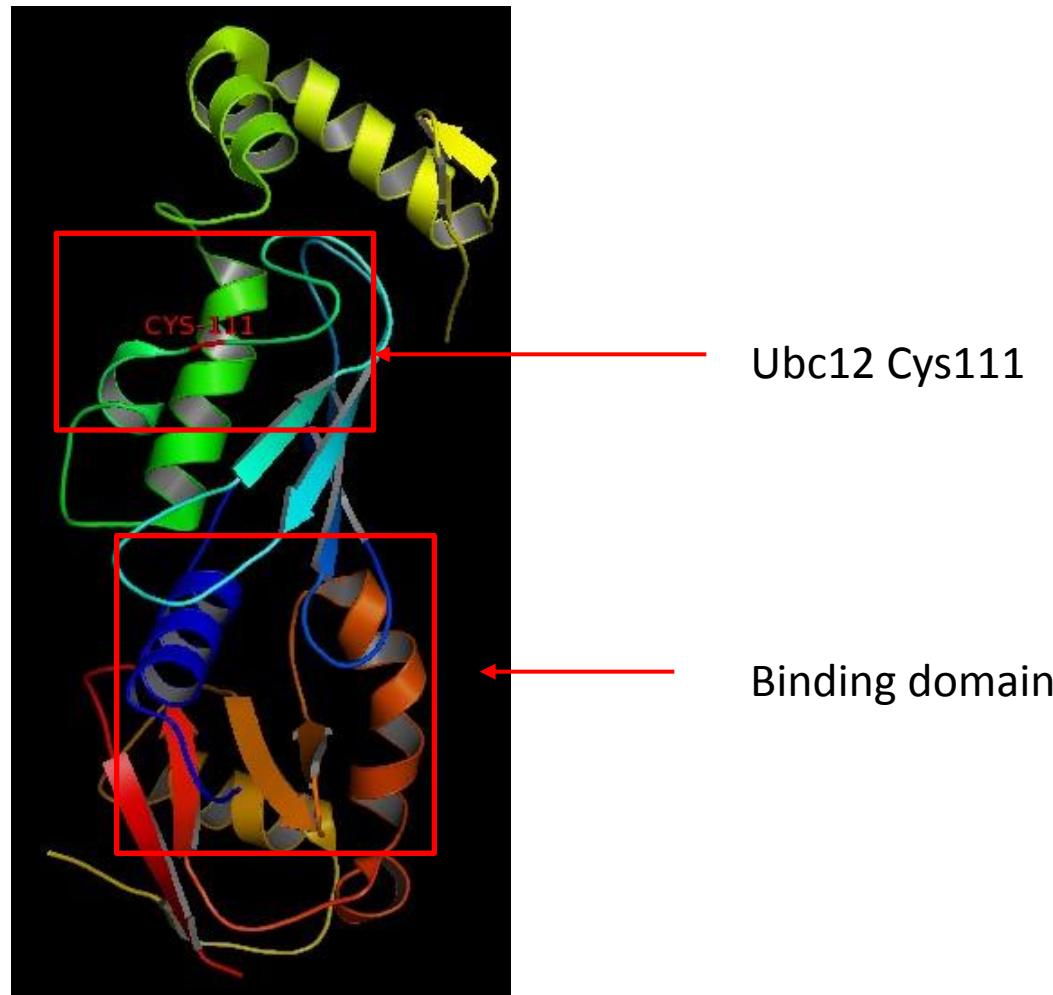
- 2. A Common Mechanism for E1-E2 Interactions



Overall structure of the Ubc12N26 complex with UBA3



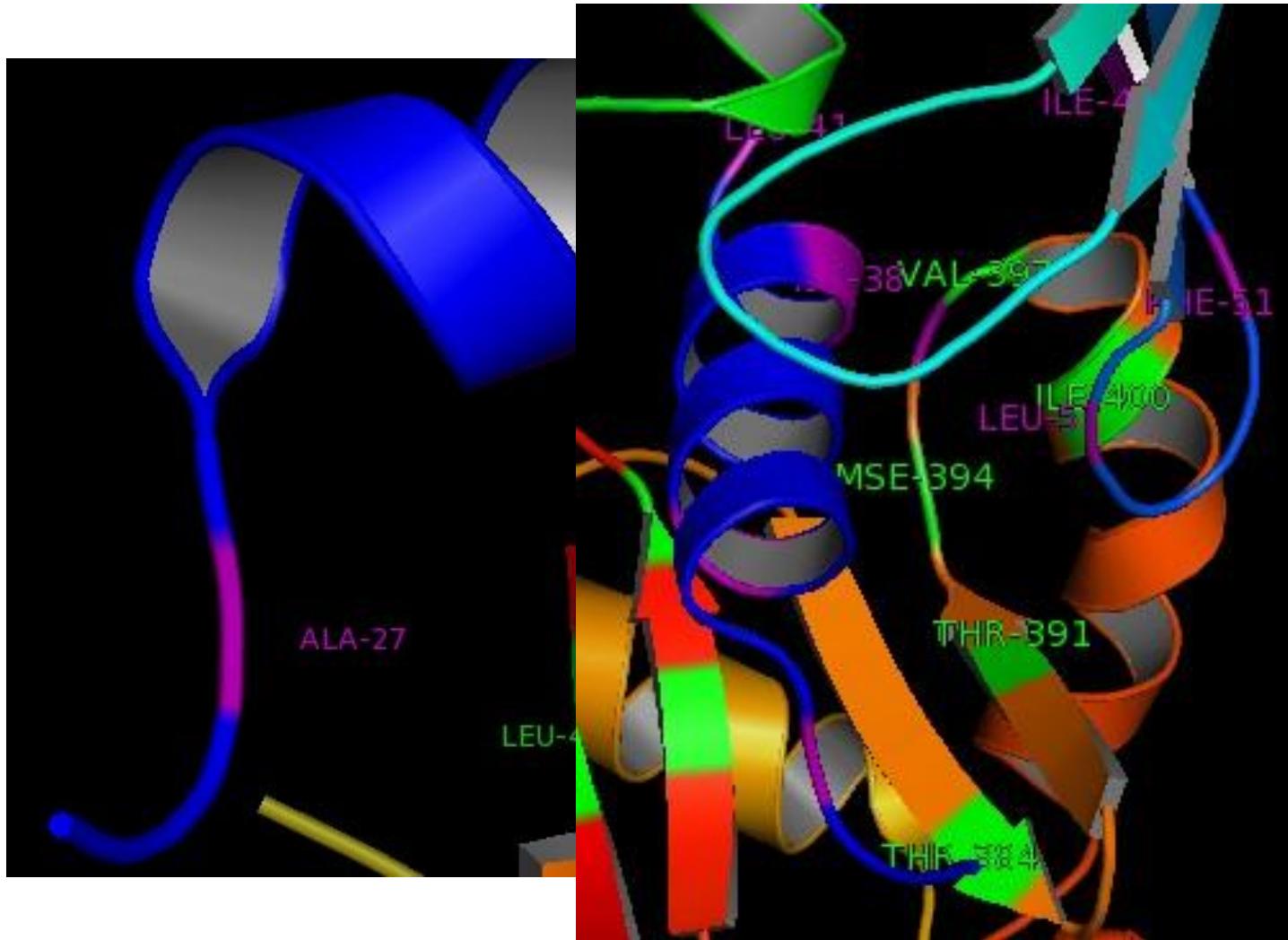
Contributions of individual residues from the Ubc12N26 to UBA3 binding



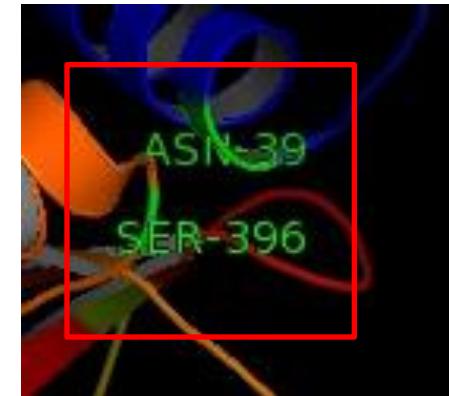
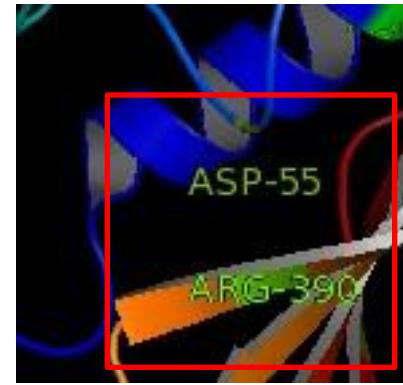
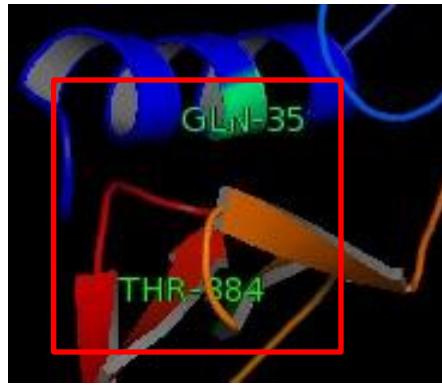
Overall Structure of the NE1^{ufd}-Ubc12^{core} Complex



Analysis



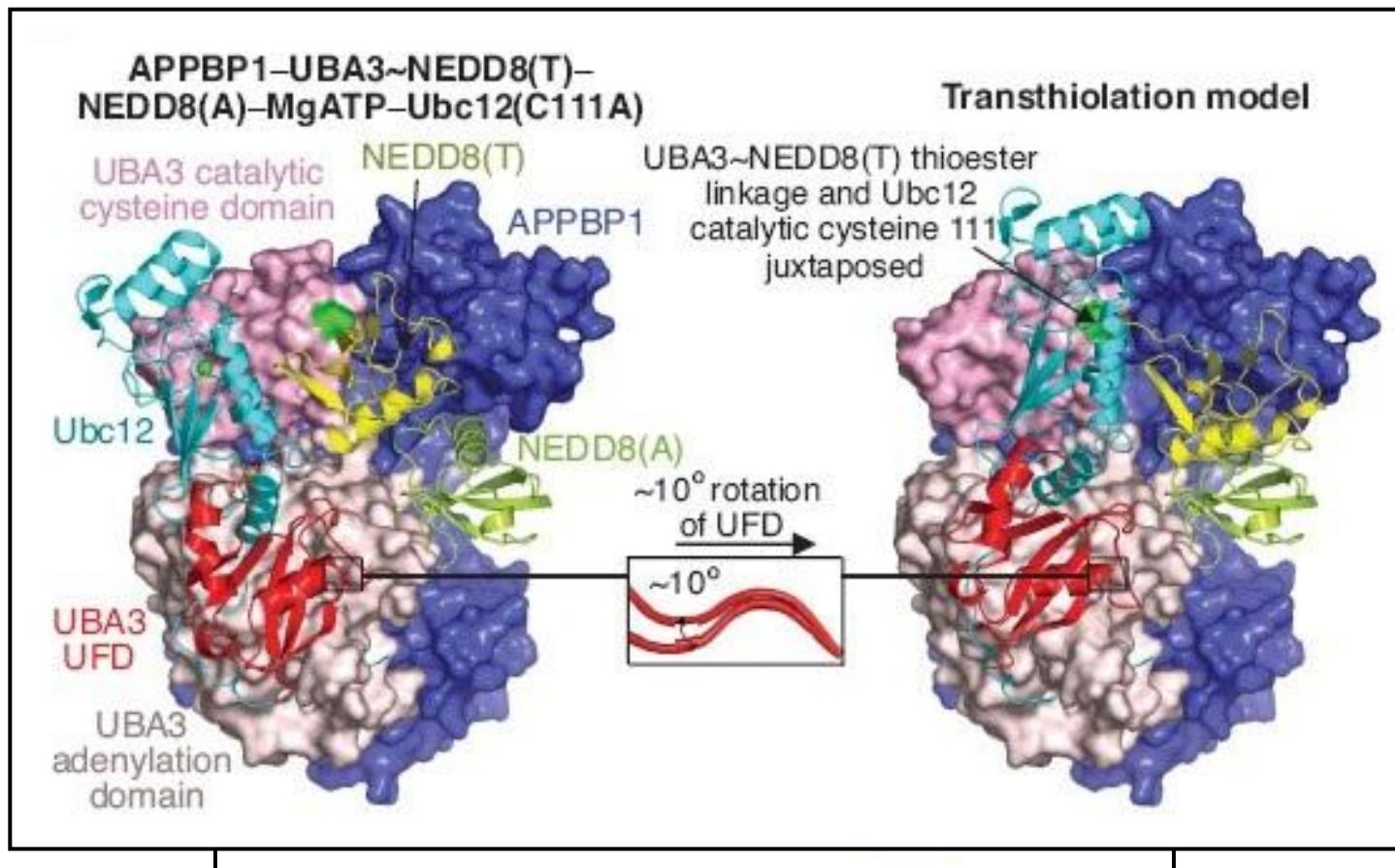
Contributions of individual residues from the Ubc12 core domain to UBA3 binding



Contributions of individual residues from the Ubc12 core domain to UBA3 binding



- 3. How does Nedd8 transfer from E1 to E2



Evidence for Rotation of the E1's Ubiquitin-Fold Domain in Ubc12 NEDD8 Thioester Formation(Danny T. Huang.et.al. 2005)

- 4. How does Nedd8 transfer from E2 to substrate
 - No Structure of Ubc12 and E3, maybe domain of E2/E1 and E2/E3 overlaps.
 - Suppose: Ubc12-Nedd8 but not free Ubc12 can interact with E3, and E3 can change its formation and then Nedd8 transfer from E2 to substrate.

Conclusion

- 1. Ubc12 is very conservative.
- 2. Both the N-terminal domain and the core domain of Ubc12 can interact with E1, this stabilize the complex between Ubc12/E1.
- 3. Transferring of Nedd8 from E1 to Ubc12 needs conformation change.
- 4. It needs more study of how Nedd8 transfers from E2 to substrate.



Thank you