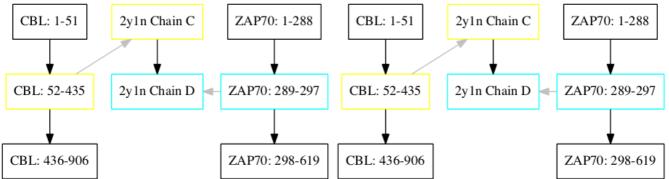
# Fusion KMT2A-CBL and Interaction 2y1n: (No direct effect on interaction 2y1n)

## **Interaction 2y1n**

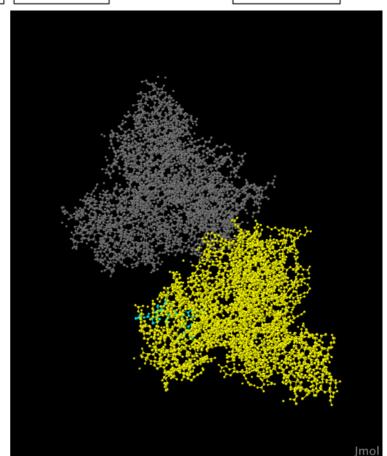
# **Fusion Effect**



### **Interactome3D**

Interaction Complex: 2y1n

- CBL
  - Chain: C
  - o Length: 1 906
  - Included in Interaction: 52 435
- ZAP70
  - o Chain: D
  - o Length: 1 619
  - o Included in Interaction: 289 297



**COSMIC** 

#### **Pfam**

- Cbl\_N: 51-175
  Cbl\_N2: 179-262
  Cbl\_N3: 264-349
  Cbl\_N3: 264-349
- Prok-RING\_4: 381-429
- UBA: 856-892

#### Mutation: COSF1971

- First Gene in Fusion: KMT2A
  - genomic breakpoint: 4086+366
  - AA breakpoint: 122
- Second Gene in Fusion: CBL
  - o genomic breakpoint: 1808-5749
  - AA breakpoint: 1916

#### **PDBe**

- Reaction: ATP + a [protein]-L-tyrosine = ADP + a [protein]-L-tyrosine phosphate.
- Author Description: Structure of c-Cbl-ZAP-70 peptide complex
- Citation: <u>Citation not listed.</u>