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## Co-IP/Pull-down In-solution Trypsin Digestion (ISD) Kit Catalog Number: MG04





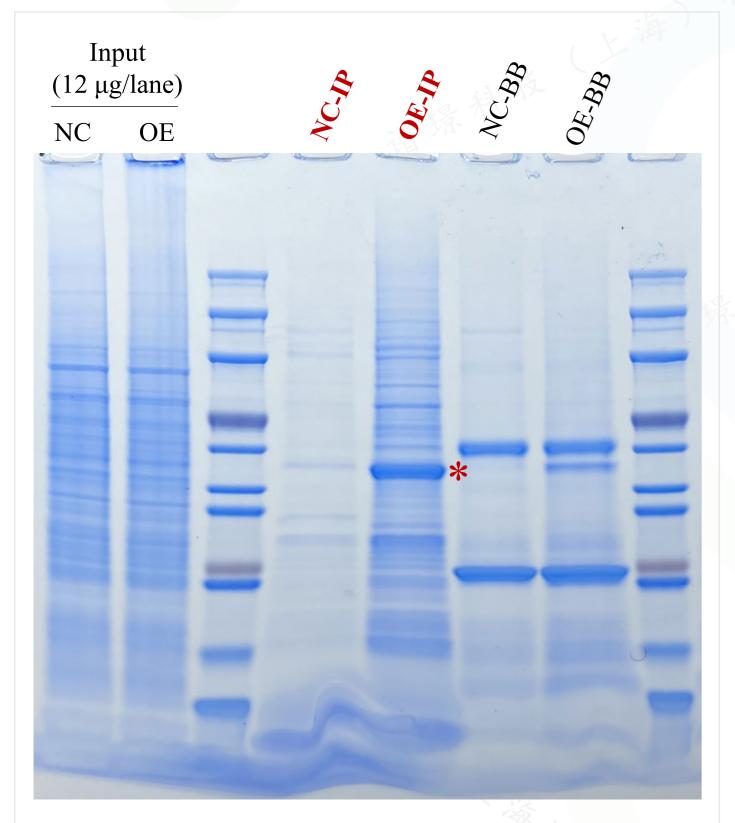
货号	名称	应用领域	质谱定量方式
MG01	IP-MS富集试剂盒 (3×FLAG tag)	蛋白质互作/后修饰筛选	LFQ [1]
MG02	SILAC-IP-MS富集试剂盒 (3×FLAG tag)	蛋白质互作/后修饰筛选	SILAC [2]
MG05	SM-PD富集试剂盒 (Biotin tag)	小分子-靶点蛋白筛选	LFQ [1]
MG08	SILAC-SM-PD富集试剂盒 (Biotin tag)	小分子-靶点蛋白筛选	SILAC [2]
MG06	SM-PD富集试剂盒 (Alkyne tag)	小分子-靶点蛋白筛选	LFQ [1]
MG09	SILAC-SM-PD富集试剂盒 (Alkyne tag)	小分子-靶点蛋白筛选	SILAC [2]
MG07	SM-PD富集试剂盒 (Azide tag)	小分子-靶点蛋白筛选	LFQ [1]
MG10	SILAC-SM-PD富集试剂盒 (Azide tag)	小分子-靶点蛋白筛选	SILAC [2]

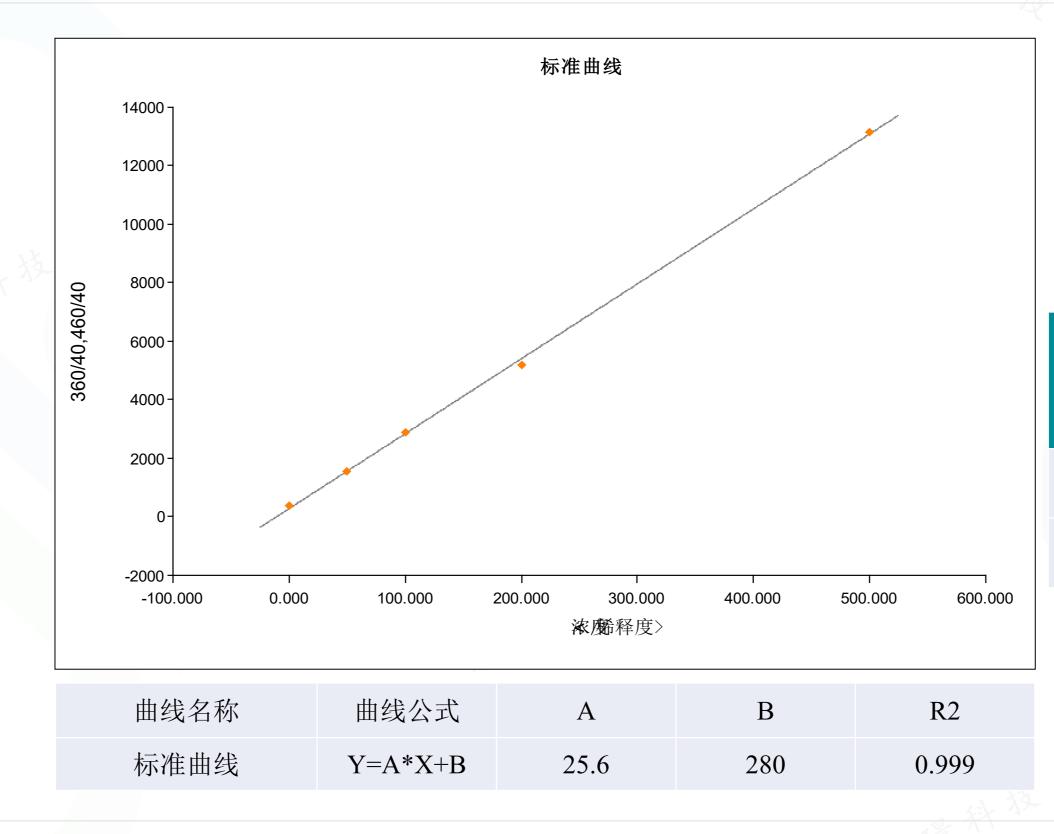
<sup>[1]</sup> Mol Cell Proteomics 2014, **13**(9):2513-2526.

<sup>[2]</sup> Mol Cell Proteomics 2002, 1(5):376-386.



## MG04 Kit: In-solution digestion of Co-IP samples prepared by MG01 Kit



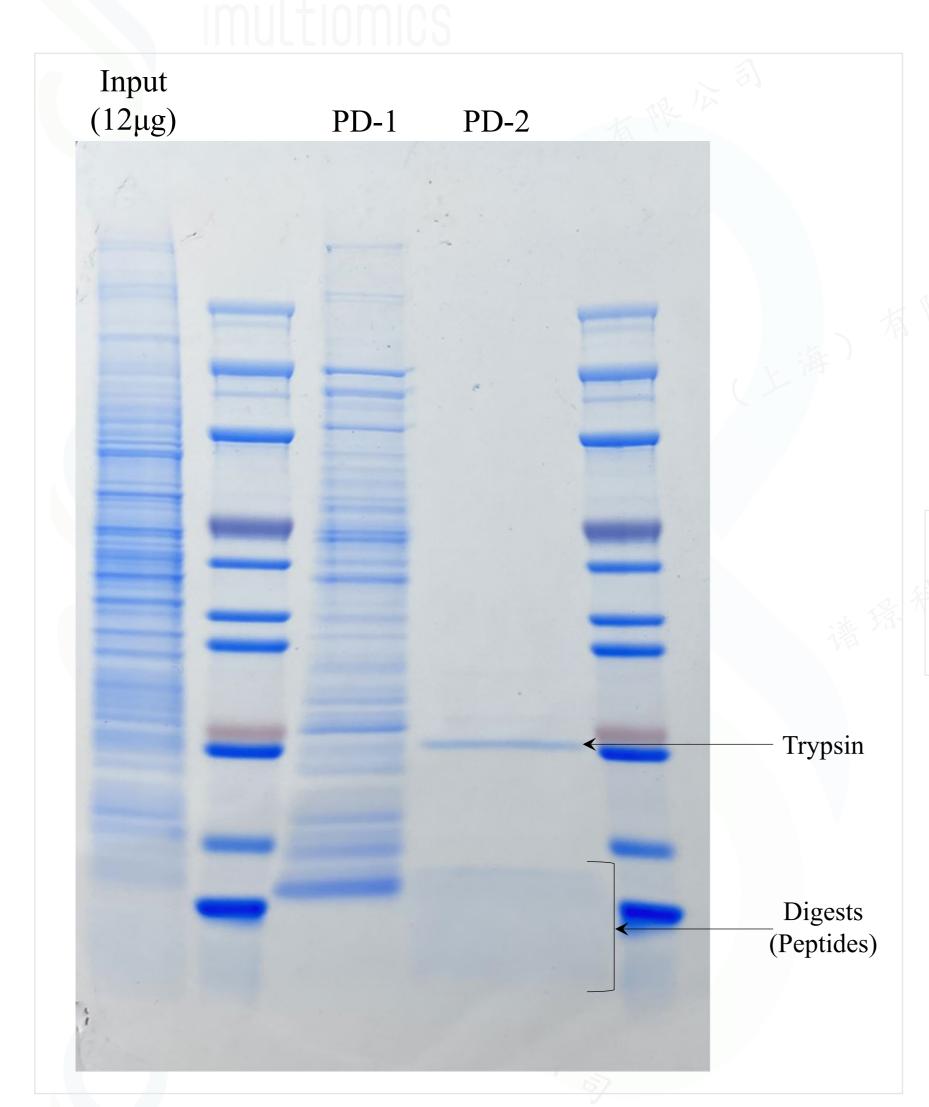


Sample	Total peptide yield (ng)
NC-IP	3437
OE-IP	8628.16

- 1. Co-IP samples of NC-IP and OE-IP were prepared according to the protocol of MG01 Kit.
- 2. (A) 70% of NC-IP and OE-IP samples was separated by SDS-PAGE gel and stained using Coomassie blue staining reagent (Imultiomics, #MGR01). Red star: Enriched 3×DYKDDDDK tag fused *Bait Protein*.
- 3. (B) 30% of NC-IP and OE-IP samples was conducted to in-solution digestion according to the protocol of MG04 Kit, and total peptide yields (ng) were measured.



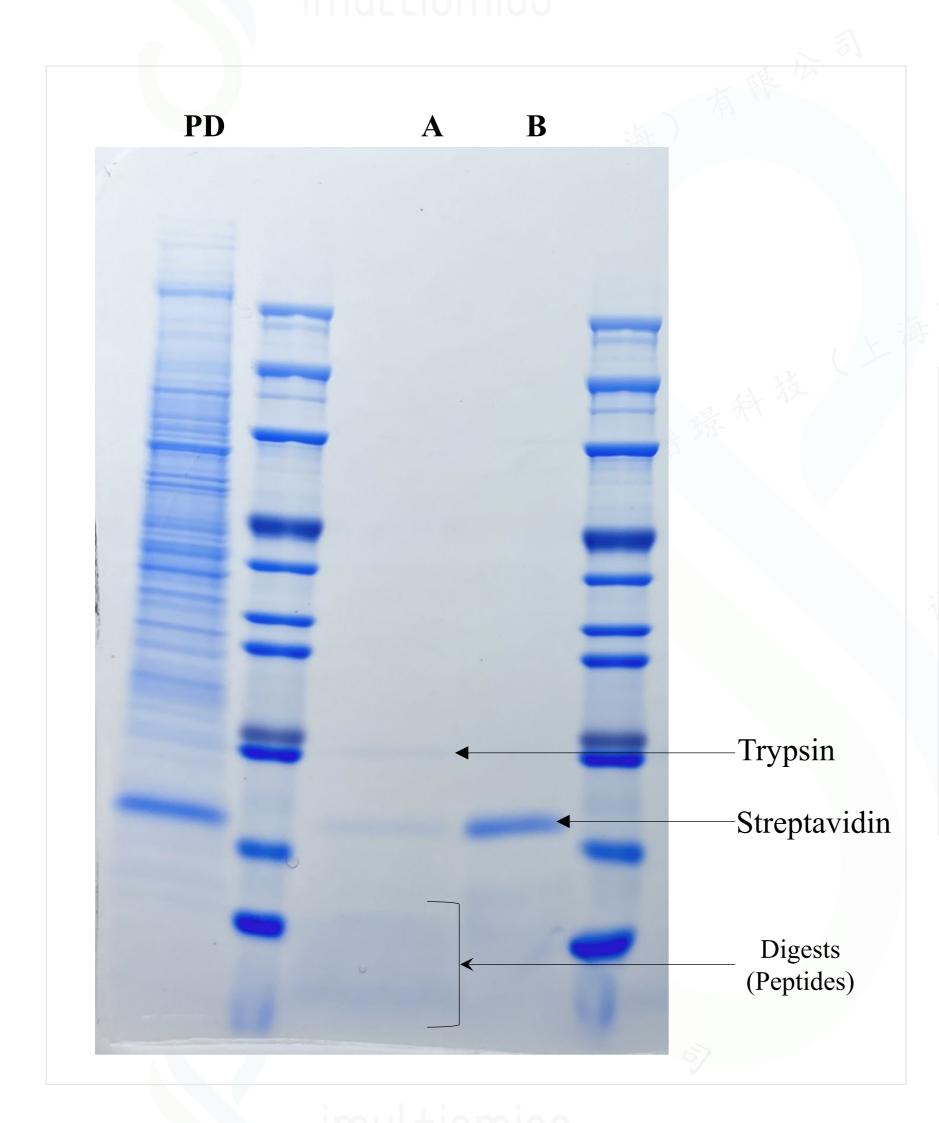
## MG04 Kit: On-beads digestion of Pull-down (PD) samples of magnetic beads



- 1. Protein pull-down (PD) was conducted using magnetic beads in duplicate (PD-1 and PD-2).
- 2. PD-2 was conducted to on-beads digestion using MG04 Kit. In contrast to PD-1, the pulled-down proteins were completely digested into peptides.
- 3. Gel was stained using Coomassie blue staining reagent (Imultiomics, #MGR01).



## MG04 Kit: On-beads digestion of Pull-down (PD) samples of agarose beads



**PD**: MG05-5 (Streptavidin-beads in MG05 Kit) enriched proteins by pull-down (PD). PD was conducted in duplicate.

**A.** PD sample was carried out on-beads digestion using MG04 Kit. the pulled-down proteins were completely digested into peptides in contrast to that of PD.

**B.** The left beads after on-beads digestion was treated with MG05-6 (5× sample loading buffer in MG05 Kit) for Coomassie blue staining reagent (Imultiomics, #MGR01). The data showed that most of streptavidin remained on beads after on-beads digestion, great for downstream LC-MS identification of pulled-down proteins.