

Evaluation of Analysis

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1 评估

筛出（瘢痕增生）能够与 TCF-AS1 结合又能与 TCF4 结合的 RNA 结合蛋白：

- 在 GEO 数据库中找到合适的数据集，与瘢痕增生有关，mRNA 和 ncRNA 的转录组数据。
- 基因共表达分析（WGCNA¹），筛选与 TCF4-AS1 lncRNA 和 TCF4mRNA 协同变化的 RNA 表达。
- 根据筛选的 RNA 寻找蛋白，获取功能注释。
- 使用预测工具²⁻⁵ 计算筛选的蛋白和 TCF4-AS1 lncRNA 和 TCF4mRNA 的结合程度。

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