## Bilinear Bandits with Low-rank Structure

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Application: Drug discovery

proteins


- Choose a pair to experiment with



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drugs


- Choose a pair to experiment with

proteins


- Goal: Find as many pairs with the desired interaction as possible

online dating

clothing recommendation


## Bilinear bandits

$$
\begin{gathered}
y=x^{\top} \Theta z+\eta \\
\text { desired interaction? (0/1) drug features protein features } \\
\text { unknown parameter }(d \text { by } d)
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- A natural model: already used for predicting drug-protein interaction.
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[Luo et al., "A network integration approach for drug-target interaction prediction and computational drug repositioning from heterogeneous information", Nature Communications, 2017]
- Issue: $d^{2}$ number of unknowns
- What if $\operatorname{rank}(\Theta) \ll d$ ?

$$
\Theta=\sum_{k=1}^{r} \sigma_{i} u_{i} v_{i}^{\top} \quad \square \sim d r \text { unknowns }
$$

- Many real-world problems exhibit the low-rank structure.


## Summary of the result

- A naïve method: reduction

$$
\mathbb{E}[y]=x^{\top} \Theta z=\left\langle\operatorname{vec}(\Theta), \operatorname{vec}\left(x z^{\top}\right)\right\rangle
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$$
\text { YES, we achieve } \frac{d^{3 / 2} \sqrt{r}}{\sqrt{T}} \quad \text { (factor } \sqrt{d / r} \text { better) }
$$

- Is this optimal?


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Let's chat.
\#127
@ Pacific Ballroom

