

# Partial Least Squares Regression (PLSR): A Detailed Breakdown

Partial Least Squares (PLS) is an extremely powerful technique—similar to PCR—but with a **major improvement**:

**It uses information from the target variable while constructing components.**

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Let's break it down clearly and thoroughly.

## What is Partial Least Squares (PLS)?

Partial Least Squares Regression (PLSR) is a supervised dimensionality-reduction + regression method used when:

- You have **many correlated predictors**
- You need to reduce dimensionality
- You want components that are **directly useful for predicting the target**

It works especially well when:

- Predictors are highly collinear
- Number of predictors  $\gg$  number of observations
- Standard linear regression becomes unstable

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PLS is most common in chemometrics, bioinformatics, spectroscopy, etc.

## How is PLS Different from PCA & PCR?

### PCA / PCR

- Finds directions of **maximum variance in  $X$  only**
- Completely ignores the target ( $y$ )
- Dimensionality reduction is unsupervised

### PLS

- Finds directions that maximize **covariance between  $X$  and  $y$**
- Actively uses the target variable while building components
- Dimensionality reduction is supervised

### Key Insight:

PCR components may not be good predictors if they have low variance but high correlation with  $y$ .  
PLS fixes this by focusing on what actually predicts the response.

## Intuition Behind PLS

Imagine you have 50 features, but only a combination of 3 of them actually matters for predicting your target.

- PCA would find components that maximize variance, not prediction strength.
- PLS finds directions in feature space that **best explain y**.

It's like asking:

"Which direction in the  $\mathbf{X}$  space helps us predict  $\mathbf{y}$  the most?"

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## How PLS Works (Step-by-Step)

Let's define:

- $\mathbf{X}$ : predictor matrix ( $n \times p$ )
- $\mathbf{y}$ : response vector ( $n \times 1$ )
- We want components  $\mathbf{t}_1, \mathbf{t}_2, \dots$

PLS builds components iteratively:

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### Step 1: Compute weights

Find weight vector  $\mathbf{w}$  that maximizes:

$$\text{cov}(\mathbf{X}\mathbf{w}, \mathbf{y})$$

This selects the direction in  $\mathbf{X}$  that has the strongest predictive relationship with  $\mathbf{y}$ .

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### Step 2: Compute the score (component)

$$\mathbf{t} = \mathbf{X}\mathbf{w}$$

This is the new transformed feature.

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### Step 3: Regress $\mathbf{y}$ on $\mathbf{t}$

$$c = \frac{\mathbf{t}^T \mathbf{y}}{\mathbf{t}^T \mathbf{t}}$$

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### Step 4: Deflate $\mathbf{X}$ and $\mathbf{y}$

Remove the information explained by the component:

$$\mathbf{X} \leftarrow \mathbf{X} - \mathbf{t}\mathbf{p}^T$$

$$\mathbf{y} \leftarrow \mathbf{y} - \mathbf{t}c$$

(where  $\mathbf{p}^T$  is the  $\mathbf{X}$  loading vector).

Now repeat for the next components.

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## The result

- You get **latent variables**  $\mathbf{t}_1, \mathbf{t}_2, \dots, \mathbf{t}_k$
  - Then you regress  $\mathbf{y}$  on these latent variables
  - Similar to PCR but the components are **optimal for prediction**
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## Why PLS Often Outperforms PCR

Because PLS uses **supervised** extraction of components, it tends to:

- ✓ Use fewer components
  - ✓ Achieve higher predictive accuracy
  - ✓ Avoid throwing away low-variance but high-predictive features
  - ✓ Handle multicollinearity better
  - ✓ Be more stable when  $p \gg n$
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## PLS vs PCR Summary Table

Table 1: Comparison of PLS and PCR

| Feature  | PCA/PCR                               | PLS                                    |
|--|---------------------------------------|--|
| Uses $\mathbf{y}$ during dimensionality reduction? | ✗ No                                  | ✓ Yes                                  |
| Maximizes  | Variance in $\mathbf{X}$              | Covariance( $\mathbf{X}, \mathbf{y}$ ) |
| Risk   | Keeps components that may not predict | Components designed to predict         |
| Performance  | Good, but sometimes unstable          | Typically better, especially with      |
| Latent features                                    | Unsupervised                          | Supervised                             |

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## When Should You Use PLS?

Use PLS when:

- You have **many correlated predictors**
- You want to reduce dimensionality **and** predict something
- You need a stable regression model
- You're analyzing chemical spectra, genetics, or high-dimensional datasets

Examples:

- NIR/IR spectroscopy analysis
  - Gene expression → disease prediction
  - High-dimensional sensor data
  - Multicollinearity-heavy datasets
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## A Simple Analogy

PCR:

“I’m going to summarize the predictors in the best way possible, even if those summaries don’t help predict  $\mathbf{y}$  very much.”

PLS:

“I’m going to summarize the predictors in the best way *for predicting  $\mathbf{y}$* , even if those summaries don’t capture all the variance.”

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