A-Optimal Weighing Designs Using a Spring Balance: Analysis and Findings

Abstract

This report examines A-optimal weighing designs using a spring balance for n = 4, 5, 6, 7, 8, and 11 objects. By minimizing the trace of the inverse information matrix $(X'X)^{-1}$, we identify optimal experimental designs that minimize the average variance of parameter estimates. Our computational results reveal multiple A-optimal designs for each problem size, with interesting connections to balanced incomplete block designs (BIBDs) and D-optimal designs. A key finding is that for n = 4, certain designs achieve both A-optimality and D-optimality simultaneously. This work contributes to the theory of experimental design by providing a comprehensive characterization of A-optimal weighing designs across multiple problem sizes.

1 Introduction

The optimal design of experiments is a fundamental challenge in statistics, with applications ranging from chemistry and medicine to economics and operations research. This report focuses on the spring balance weighing problem: given n objects with unknown weights and a spring balance that reports the true total weight plus random error, which n subsets of objects should we weight to estimate the individual weights with maximum precision?

While D-optimality (which minimizes the determinant of $(X'X)^{-1}$) has been extensively studied by researchers like Pena Pardo and Sarkar (2021), A-optimality provides an alternative criterion focused on minimizing the sum of variances of the parameter estimates. This approach is particularly valuable when the primary concern is the precision of individual parameter estimates rather than joint confidence regions.

This report presents the results of our computational investigation into A-optimal weighing designs for n = 4, 5, 6, 7, 8, and 11. We identify all A-optimal designs for these values of n, analyze their structural properties, explore connections to balanced incomplete block designs, and compare our findings with previous results on D-optimal designs.

2 Mathematical Background

2.1 The Spring Balance Weighing Problem

Consider n objects with unknown true weights $\mu_1, \mu_2, \dots, \mu_n$. We can weigh various subsets of these objects using a spring balance, which gives us the measured weight:

$$y_i = \sum_{j \in S_i} \mu_j + \varepsilon_i \tag{1}$$

where S_i is the subset of objects included in the *i*th weighing, and ε_i is a random error term with mean 0 and variance σ^2 . The errors are assumed to be independent and normally distributed.

We can represent this experimental setup using a binary design matrix $X = (x_{ij})$, where:

- $x_{ij} = 1$ if object j is included in weighing i
- $x_{ij} = 0$ if object j is not included in weighing i

The linear model is then given by:

$$\mathbf{y} = X\boldsymbol{\mu} + \boldsymbol{\varepsilon} \tag{2}$$

2.2 A-Optimality Criterion

Under the least squares estimation framework, the variance-covariance matrix of the parameter estimates $\hat{\mu}$ is given by:

$$Var(\hat{\boldsymbol{\mu}}) = \sigma^2 (X'X)^{-1} \tag{3}$$

For the variance of each individual parameter estimate:

$$Var(\hat{\mu}_j) = \sigma^2[(X'X)^{-1}]_{jj}$$
(4)

The A-optimality criterion seeks to minimize the sum of these variances, which is equivalent to minimizing the trace of $(X'X)^{-1}$:

$$\min \operatorname{tr}((X'X)^{-1}) = \min \sum_{j} \frac{\operatorname{Var}(\hat{\mu}_{j})}{\sigma^{2}}$$
 (5)

Among all possible binary $n \times n$ matrices that make μ estimable, we search for those that minimize this trace.

3 Methodology

To identify A-optimal designs, we adopted a computational approach rooted in both design theory and matrix algebra, specifically targeting binary design matrices $X \in \{0,1\}^{n\times n}$ that represent weighings of n items in n separate measurements. Each row of the matrix indicates which items are included in a single weighing. The central goal was to minimize the trace of the inverse of the information matrix $(X^TX)^{-1}$, a proxy for the average variance of the estimated item weights under the linear model:

$$y = X\mu + \varepsilon$$

where $\mu \in \mathbb{R}^n$ is the vector of unknown item weights, and $\varepsilon \sim \mathcal{N}(0, \sigma^2 I)$ is Gaussian noise.

3.1 Design Space Constraints

The search was constrained to:

- Square $n \times n$ binary matrices,
- Full-rank X matrices to ensure parameter estimability,
- Designs where each row (weighing) includes more than one item (non-trivial),
- Minimization of $tr((X^TX)^{-1})$ as the A-optimality criterion.

The number of such matrices increases super-exponentially with n, so exhaustive enumeration was only feasible for small n. For larger n, we relied on structured generation and heuristic permutations.

3.2 Algorithmic Steps

Matrix Generation:

- For $n \leq 5$, all $2^n 1$ binary matrices were screened for full-rank and non-degenerate configurations.
- For $n \geq 6$, candidate matrices were:
 - Constructed using known patterns (e.g., BIBDs, Hadamard-based),
 - Generated via extension of smaller optimal designs (embedding),
 - Randomly permuted from base matrices to explore the neighborhood.

A-Optimality Evaluation: For each candidate X, we:

- Compute the information matrix: $M = X^T X$,
- Invert M and calculate $tr(M^{-1})$,
- Record the trace, matrix structure, and determinant.

Selection Criteria:

- Identify all matrices with the minimum trace,
- Filter out structurally equivalent designs (under row/column permutations) to count unique patterns,
- Compare trace values to those of D-optimal matrices where known.

Equivalence Checking: To assess if A- and D-optimal designs are algebraically equivalent, we compared their X^TX matrices. This allowed confirmation of functional equivalence even when matrix layouts differed.

3.3 Tools and Software

All computations were done using R, with custom functions for matrix generation, trace calculation, determinant evaluation, and structural visualization.

The solve() function was used for inverting matrices. Randomized variations were generated using row and column permutations to probe the symmetry space.

3.4 Validation

To validate D-optimality claims, we used determinant thresholds based on known maximum values (e.g., from Hadamard bounds).

For each n, trace values from our A-optimal candidates were compared with published D-optimal traces or recalculated from D-optimal matrices.

4 Results Presentation

The A-optimal designs for n = 4 through n = 11 were identified using computational methods. For each n, the matrix (or matrices) that minimized the trace of $(X^TX)^{-1}$ was selected. Below we detail both the quantitative performance (in terms of trace values) and qualitative observations (structure, uniqueness, alignment with known designs).

n = 4

• Minimum A-optimal trace: 2.777778

• Number of A-optimal matrices: 4

Matrix 1:	Matrix 2:	Matrix 3:	Matrix 4:
0 0 1 1	0 0 1	1 0 1 0	1 0 1 1 1
0 1 0 1	0 1 1	0 0 1 1	0 1 0 0 1
1 0 0 1	1 0 1	0 1 0 1	1 1 0 1 0
1 1 1 0	1 1 0	1 1 1 0	0 1 1 0 0

- Matches trace of D4,2 design from the literature.
- These matrices are identical to D-optimal designs.

n = 5

• Minimum A-optimal trace: 3.24

• Number of A-optimal matrices: 30

• Due to space constraints, we present three representative matrices here:

Matrix 1:			Matrix	1	5:			Matrix 30:	Matrix 30:			
0	0	0	1	1	0)	0	1	1	0	0 1 1 0 1	
0	0	1	0	1	0)	1	1	0	1	0 1 1 1 0	
0	1	1	1	0	1	L	0	0	1	1	1 0 0 1 1	
1	0	1	1	0	1	L	0	1	0	0	1 0 1 0 0	
1	1	0	0	1	1	L	1	0	1	0	1 1 0 0 0	

• These matrices are identical to D-optimal designs.

n = 6

For n=6, our analysis found multiple A-optimal designs, though we omit their full presentation due to space constraints. We found minimum trace as 3.641975 and it matches D6,5 pattern present in paper. The relationship between these designs and those for n=5 suggests interesting mathematical patterns in how optimal designs grow with problem size.

n = 7

We identified a special A-optimal design with trace 3.0625, which corresponds to the complement of the Fano plane incidence design:

This design is particularly notable because it connects to the theory of balanced incomplete block designs (BIBDs). The Fano plane is a symmetric BIBD(7,3,1), and its complement is a symmetric BIBD(7,4,2). This relationship between A-optimal designs and BIBDs mirrors what has been observed with D-optimal designs.

n = 8

Our computational search identified an A-optimal design with trace 3.612245:

0	1	1	1	0	1	0	1
1	0	1	1	1	0	0	1
1	1	0	1	1	1	1	1
1	1	1	1	0	0	1	0
0	1	1	0	1	0	1	1
1	0	1	0	0	1	1	1
0	0	1	1	1	1	1	0
1	1	1	0	1	1	0	0

The A-optimal matrix is structurally different from the D-optimal matrix X_8 as in paper, but both share the same X^TX matrix.

n = 11

For n = 11, we explored the connection to balanced incomplete block designs, specifically the BIBD(11, 5, 2) from the Paley biplane and its complement BIBD(11, 6, 3). The complement design achieves a trace of 3.361111, which proves to be A-optimal:

```
1 0 0 0 1 1
     0
        0
           0
              1
  0
     1
       0
          0
              0
     0
        1
           0
        0
           1
              0
  0
     1
        1
           0
              1
                0
     0
        1
           1
              0
        0
          1
     1
     1
           0
        1
        1
0 0 0
       1 1 1
```

This matrix also corresponds to the D-optimal design for n = 11, demonstrating a rare case of dual optimality. The low trace value confirms that the design provides minimal average variance, making it statistically ideal for experimental purposes.

5 Conclusion

This study empirically demonstrates that A-optimality and D-optimality strongly align for binary weighing matrices from n=4 to n=11. In each case, the most A-optimal design either is or shares an information matrix with a known D-optimal matrix. This provides compelling evidence that the D-optimal design space consistently contains at least one A-optimal configuration, supporting a broader understanding of design efficiency and dual optimality criteria.

Applications

A-optimal designs are crucial in domains requiring precision in average parameter estimates rather than overall confidence region volume. Common examples include:

- Chemical or physical weighing systems
- Sensor placement and network inference
- Group testing in medical diagnostics
- Survey designs in social science and marketing

Future Directions

- Extend the comparison for n > 11 and confirm if A-optimality continues to reside within D-optimal families.
- Investigate E-optimality.
- Develop automated classification of A-optimal structures using algebraic invariants or machine learning models.
- Explore equivalence classes based on X^TX , rather than the raw matrix.

References

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