

Supplementary Materials for
**Precursor recommendation for inorganic synthesis by machine learning
materials similarity from scientific literature**

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Supplementary Text

Model training and validation

To train the PrecursorSelector encoding model, 44,736/2,254/2,934 synthesis reactions were used as the training/validation/test set as discussed in Section “Data preparation”. Each reaction consists of a target material and precursor materials as extracted from the literature. For the purpose of training and validation, a random subset of precursor materials is selected to be replaced with a placeholder “[MASK]” (21) in each reaction, referred to as the masked reaction. Because a combinatorial number of masked reactions can be generated from the same reaction, the sampling space of masked reactions is much larger than that of original reactions. To sample as many different masked reactions during the training phase, we employ a dynamic masking strategy (74) that randomly samples a batch of reactions and re-generates the masking pattern in every training step. Different from the training samples, the validation samples are generated using static masking during data preprocessing because keeping the validation set unchanged is necessary for model selection afterwards. In this work, we trained with a batch size of 8 masked reactions for 500,000 steps, or 50 epochs with 10,000 steps per epoch. The optimizer used was Adam (75) with learning rate of 5×10^{-4} , $\beta_1 = 0.9$, and $\beta_2 = 0.999$. Starting from the 2,254 original reactions in the validation set, we applied the masking procedure and randomly sampled 3,320 masked reactions for validation. The optimal model selected was the one with minimal loss on the validation samples to minimize overfitting (76).

Two tasks are implemented in the representation learning model: (1) the masked precursor completion (MPC) task that predicts the complete precursor set based on the target material and the synthesis context provided by the unmasked precursors, and (2) the composition recovery task that predicts the chemical composition of the target material from the encoded target vector. The loss function in the MPC task, denoted as L_1 , is the circle loss (71) to maximize the within-class similarity and minimize the between-class similarity in multi-label classification. Here, the within-class similarity corresponds to the similarity of precursor materials present in the same reaction, while the between-class similarity corresponds to the similarity between used and unused precursor materials. The loss function in the composition recovery task, denoted as L_2 , is the mean squared error (MSE) loss to compare the difference between predicted composition and the real composition of the target material. The total loss, denoted as L_{multi} , is an adaptive multi-task loss (73) to automatically weigh L_1 and L_2 with

$$L_{multi} = \frac{1}{\sigma_1^2} L_1 + \frac{1}{\sigma_2^2} L_2 + \log \sigma_1^2 + \log \sigma_2^2 \quad (S1)$$

where σ_1 and σ_2 are the model’s observation noise parameters which are learned alongside other model parameters. The training and validation losses for each task and the total are shown in Fig. S1. The training loss is averaged every 500 training steps to estimate performance on a substantial number of training samples, which in this study is 4,000. The validation loss is evaluated before training and at the end of each training epoch. As the training loss continues to decrease, the validation loss initially decreases and then increases. The minimal total validation loss is achieved at the end of 15th epoch, leading to the optimal model. The final performance of the optimal model is tested by predicting precursors for 2,654 unseen target materials in the test set. Our similarity-based recommendation strategy successfully reproduces a known precursor set 82% of the time in five attempts or less.

Computation time for similarity evaluation

In this work, all 24,034 materials in the knowledge base are converted to 32-dimensional vectors in advance, forming a $24,034 \times 32$ matrix. For the 2,654 test materials, we monitored the time required to vectorize them one by one and calculate their cosine similarity to the target vectors in the pre-stored matrix. The similarity evaluation took merely 26 seconds for all the test materials (i.e. 0.01 seconds/material) because of the fast matrix multiplication.

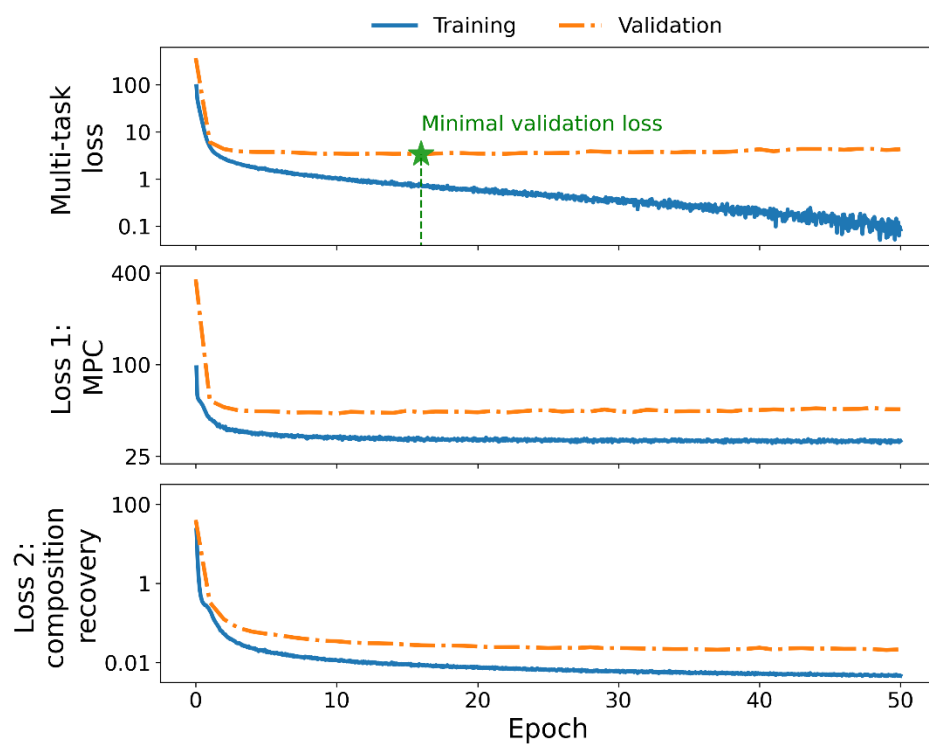


Fig. S1. Evolution of training and validation loss while training the PrecursorSelector encoding model. Top: total multi-task loss, L_{multi} . Middle: loss for the MPC task, L_1 . Bottom: loss for the composition recovery task, L_2 .

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