

Screening

In **Screening**, you select which studies to include in your nest and which should be excluded (with reasons).



Video

Steps to Screen a Nest:

1. Configure Screening

Before screening individual studies, you must [Configure Screening](#). To Configure exclusion reasons, list all reasons that a record would be excluded from your nest.

2. Select Screening Mode

There are several Screening Modes that you can select to alter the flow of records through your nest. Each option can be toggled on the Admin page for your nest:

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Screening Mode

In Standard Screening, one user screens each record. Inclusion sends the record forward for gathering, such as tagging, extraction, and Risk of Bias assessment. Exclusion does not queue the record for gathering. This determination may be updated at any time.

In Dual Screening, two users independently screen each record, and then all screening determinations are reviewed by an administrator. The administrator adjudicates any disagreement between the original screeners and sets the final determination for each record.

In Two Pass Screening, all records are first rapidly screened using only title and abstract. Records may be advanced from title/abstract screening to more intensive full text screening, where final inclusion is determined.

Standard Dual Two Pass

Screening

Inclusion Modeling

Inclusion models predict the probability of individual records being included during screening, using your past screening decisions. These probabilities help AutoLit determine which studies to show first during the screening process to get you screening faster.

The model can be trained manually or automatically (recommended). If the inclusion model is set to automatic, the model will be retrained after every 10 newly screened records. Otherwise, the model can be trained and retrained manually during screening. Your nest must contain at least 1 inclusion and 10 records in order to train a model.

☐ Automatic Training View Inclusion Model

Hiding the Model

Probabilities predicted by the model may be displayed during screening to speed up work or hidden if you wish to minimize bias. Studies will still be ordered by inclusion probability, even when hidden. To completely remove probabilities and ordering, delete the existing inclusion model and turn off automatic training.

☐ Hide Probabilities

Extraction

Screening Modes:

- **Standard Screening:** Each record must be Included or Excluded by a single user, with the

option of marking Full Text Review.

- **Dual Screening:** Each record must be Included or Excluded by two independent users, and then an Admin must adjudicate all disagreements. Each user and Admin has the option of marking Full Text Review.
- **Robot Screening:** Once enough records are screened to train the Inclusion Probability Model with enough accuracy, each record must be Included or Excluded by one reviewer, and one 'robot'. Then, the Admin must adjudicate all disagreements. Each user and Admin has the option of marking Full Text Review.
- **Two-Pass Screening:** Each record must be Advanced or Excluded at the Abstract level by one user, and Included or Excluded based on a second Full Text Review step.
- **Dual Two-Pass Screening:** Each record must be Advanced or Excluded at the Abstract level by *two* users, and Included or Excluded based on *two* reviewers' designations during a second Full Text Review step. All disagreements at each step must be adjudicated by an Admin.

The best practice is to **choose a Screening Mode before you begin Screening**. While you can switch between Screening Modes at any point, and final Inclusion/Exclusion decisions can be carried over, switching modes after beginning Screening **can result in lost data**.

3. Configure Your Keywords

To aid users in screening, you can [configure keywords](#) either in the Configure Screening page or on the Abstract view of any study in your nest. These User Keywords will be underlined in all abstracts.

These are different from the P, I, and O highlighting in that the highlighting is automatically generated by our RoboPICO software based on the abstract's contents, while Your Keywords are custom-added as described here.

4a. Include and Exclude Records

The core activity of Screening is [Including and Excluding](#) records. In this process, you review the abstracts of records and exclude any that meet your Exclusion Reasons. For any article that is includable (as in, no Exclusion Reason applies), select "Include" and (optionally) upload the full text. There is also an option to "unscreen" a study. You can do this on [individual studies](#) or across many studies in a [bulk action](#).

Uploading Full Texts

Generally, you should complete [Full Text PDF Upload](#) for any record that you Include, so that it is available to Tag during the next stage.

The exception: If you set up the **Two-Pass Screening Mode** (see above), you should upload full texts for all studies that are Advanced to Full Text Review.

4b. Related reports

While screening, it isn't unusual to find the same data reported across multiple papers, and you will

most likely include both the primary papers and all the secondary analyses. For the non-primary papers with data from the same trial, you can mark them as [related reports](#) to keep track of them and to have the PRISMA adjusted to reflect the difference in the number of studies versus the number of reports.

5. When are you finished Screening?

You are finished Screening when all records in your nest are either Included or Excluded, as indicated by the Screening progress bar. At this point, you should move on to [Tagging](#) and [Extracting](#) the included records!

Additional Tools

- [Screening Model](#): How to use the built-in Inclusion Prediction to assist in ordering records and informing your screening.
- [Add Exclusion Reasons On the Fly](#): How to add Exclusion Reasons while you are screening without leaving the page.
- [Exclusion Reason Sets](#): How to import Organizational Templates containing pre-configured Sets of Exclusion Reasons.

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