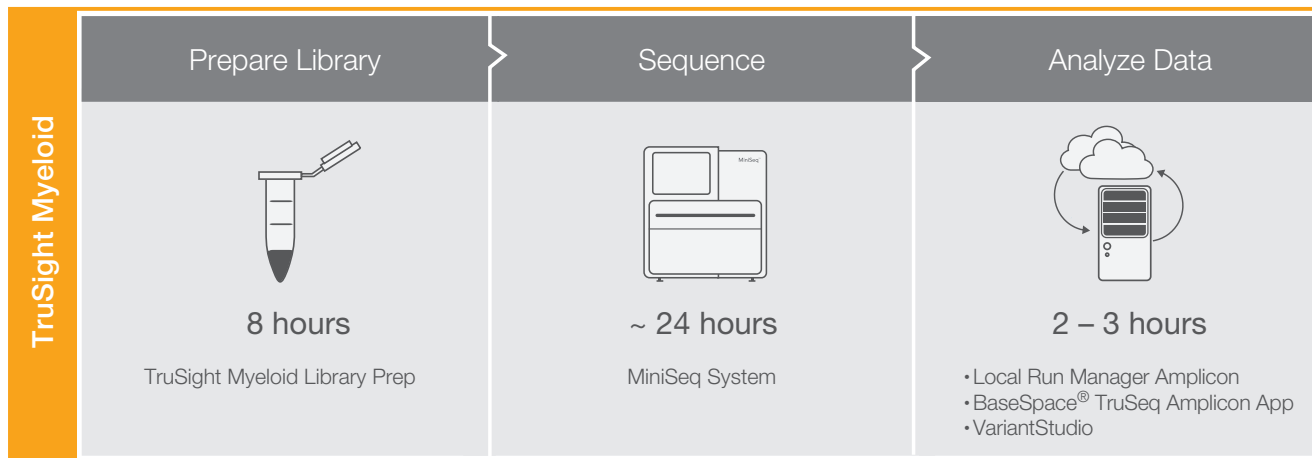




# TruSight<sup>®</sup> Myeloid Workflow on the MiniSeq<sup>™</sup> System



This checklist supports the following workflow choices.

Sequencing Instrument:	MiniSeq System
Setup Option	Local Run Manager
Library Preparation Kit:	TruSight Myeloid Sequencing Panel
Indexing:	Dual Indexing
Reagent Kits:	MiniSeq Kit
Analysis Workflow:	Amplicon
Analysis Software:	Local Run Manager

MiniSeq sequencing using TruSight Myeloid Sequencing Panel libraries

## Set Run Parameters

- 1 Log in to Local Run Manager.
- 2 Click **Create Run**, and select **Amplicon**.
- 3 Enter a run name that identifies the run.
- 4 [Optional] Enter a run description.
- 5 From the Library Kit drop-down list, select TruSight Amplicon Panels.
- 6 Specify the number of cycles for the run.
- 7 Select a variant calling method.
- 8 Click **Show advanced module settings** and specify the Read Stitching and Variant Quality Filter settings.
- 9 Click **Import Manifests**.
- 10 Navigate to the manifest file.
- 11 Enter a unique sample ID.
- 12 [Optional] Enter a sample description.
- 13 Select an Index 1 adapter.
- 14 Select an Index 2 adapter.
- 15 Select a manifest file.
- 16 Select a reference genome.
- 17 Click **Save Run**.

## Hybridize Oligo Pool

- 1 Add 5  $\mu$ l ACD1 and 5  $\mu$ l TE or water to 1 well of the HYP plate.
- 2 Add 10  $\mu$ l gDNA to each remaining well.
- 3 Add 5  $\mu$ l TSO to each well containing gDNA.
- 4 Centrifuge at 1000  $\times$  g for 1 minute.
- 5 Add 35  $\mu$ l OHS2. Pipette to mix.
- 6 Centrifuge at 1000  $\times$  g for 1 minute.
- 7 Place on the preheated heat block and incubate for 1 minute.
- 8 Reset the temperature to 40°C and incubate for 80 minutes.

## Remove Unbound Oligos

- 1 Make sure that the heat block has cooled to 40°C.
- 2 Remove from the heat block.
- 3 Centrifuge at 1000 × g for 1 minute.
- 4 Transfer each sample to the FPU plate.
- 5 Cover and centrifuge at 2400 × g for 5 minutes.
- 6 Wash 2 times with 45 µl SW1.
- 7 Reassemble the FPU plate.
- 8 Add 45 µl UB1.
- 9 Cover and centrifuge at 2400 × g for 5 minutes.

## Extend and Ligate Bound Oligos

- 1 Add 45 µl ELM4 to the FPU plate.
- 2 Incubate at 37°C for 45 minutes.

## Amplify Libraries

- 1 Arrange the Index 1 (i7) adapters in columns 1–12.
- 2 Arrange the Index 2 (i5) adapters in rows A–H.
- 3 Place the IAP plate on a TruSeq Index Plate Fixture.
- 4 Use a multichannel pipette to add 4 µl of each Index 1 (i7) adapter to each row.
- 5 Use a multichannel pipette to add 4 µl of each Index 2 (i5) adapter to each column.
- 6 Add 56 µl TDP1 to 2.8 ml PMM2.
- 7 Invert to mix.
- 8 When incubation is complete, remove the FPU plate from the incubator and remove the seal.
- 9 Cover and centrifuge at 2400 × g for 5 minutes.
- 10 Use a multichannel pipette to add 25 µl 50 mM NaOH.
- 11 Incubate at room temperature for 5 minutes.
- 12 Transfer 22 µl PMM2/TDP1 master mix to the IAP plate.
- 13 Transfer samples eluted from the FPU plate to the IAP plate.
- 14 Centrifuge at 1000 × g for 1 minute.
- 15 Transfer to the post-amplification area.
- 16 Determine the required number (X) of PCR cycles using the following table:

Number of Amplicons in TSO	Number of PCR Cycles (X)
< 96 amplicons	33
97–384 amplicons	28
385–768 amplicons	27
769–1536 amplicons	26

- 17 Perform PCR on a thermal cycler using the following program:

- ▶ 95°C for 3 minutes
- ▶ X cycles of:
  - ▶ 95°C for 30 seconds
  - ▶ 66°C for 30 seconds
  - ▶ 72°C for 60 seconds
- ▶ 72°C for 5 minutes
- ▶ Hold at 10°C

**SAFE STOPPING POINT**

If you are stopping, leave the plate on the thermal cycler at 2°C to 8°C overnight.

**Clean Up Libraries**

- 1 Centrifuge the IAP plate at 1000 × g for 1 minute.
- 2 Run an aliquot of the libraries on 4% agarose gel (5 µl) or Bioanalyzer (1 µl).
- 3 Add 45 µl AMPure XP beads to the CLP plate.
- 4 Transfer all the supernatant from the IAP plate to the CLP plate.
- 5 Shake at 1800 rpm for 2 minutes.
- 6 Incubate at room temperature for 10 minutes.
- 7 Place on a magnetic stand until liquid is clear.
- 8 Remove and discard all supernatant.
- 9 Wash 2 times with 200 µl 80% EtOH.
- 10 Use a 20 µl pipette to remove residual EtOH.
- 11 Remove from the magnetic stand and air-dry for 10 minutes.
- 12 Add 30 µl EBT.
- 13 Shake at 1800 rpm for 2 minutes.
- 14 Incubate at room temperature for 2 minutes.
- 15 Place on a magnetic stand until liquid is clear.
- 16 Transfer 20 µl supernatant from the CLP plate to the LNP plate.
- 17 Centrifuge at 1000 × g for 1 minute.

**SAFE STOPPING POINT**

If you are stopping, seal the plate and store at 2°C to 8°C for up to 3 days. Alternatively, store at -25°C to -15°C for up to 7 days.

**Normalize Libraries**

- 1 Add 4.4 µl LNA1 per library to a new 15 ml conical tube.
- 2 Use a P1000 pipette to resuspend LNB1.
- 3 Transfer 800 µl LNB1 to the tube of LNA1.
- 4 Add the LNA1/LNB1 mix to a trough.
- 5 Add 45 µl LNA1/LNB1 to the LNP plate.
- 6 Shake at 1800 rpm for 30 minutes.
- 7 Place on a magnetic stand until liquid is clear.
- 8 Remove and discard all supernatant.
- 9 Remove from the magnetic stand.
- 10 Wash 2 times with 45 µl LNW1.
- 11 Remove residual LNW1.
- 12 Remove from the magnetic stand.
- 13 Add 30 µl fresh 0.1 N NaOH.
- 14 Shake at 1800 rpm for 5 minutes.
- 15 Place the LNP plate on a magnetic stand until liquid is clear.
- 16 Add 30 µl LNS2 to the SGP plate.
- 17 Transfer 30 µl supernatant from the LNP plate to the SGP plate.
- 18 Centrifuge at 1000 × g for 1 minute.

**SAFE STOPPING POINT**

If you are stopping, seal the plate and store at -25°C to -15°C for up to 30 days.

## Pool Libraries

- 1 Transfer 5 µl to an 8-tube strip.
- 2 Seal the plate and store at -25°C to -15°C.
- 3 Transfer the contents of the 8-tube strip to the PAL tube.

## Prepare Consumables

- 1 Remove the reagent cartridge from -25°C to -15°C storage.
- 2 Thaw reagents in a room temperature water bath for 90 minutes.
- 3 Invert the cartridge 5 times to mix reagents.
- 4 Gently tap on the bench to reduce air bubbles.
- 5 Remove a new flow cell package from 2°C to 8°C storage.
- 6 Set the unopened flow cell package aside at room temperature for 30 minutes.
- 7 Remove the flow cell from the foil package and flow cell container.
- 8 Clean the glass surface of the flow cell with a lint-free alcohol wipe.
- 9 Dry with a lint-free lens cleaning tissue.



## Denature, Dilute, and Load Libraries

- 1 Thaw the Hybridization Buffer at room temperature.
- 2 Vortex briefly before use.
- 3 Preheat the incubator to 98°C.
- 4 Combine the 5 µl pooled libraries and 995 µl prechilled Hybridization Buffer in a microcentrifuge tube.
- 5 Vortex briefly and then centrifuge at 280 × g for 1 minute.
- 6 Transfer 250 µl diluted library to a new microcentrifuge tube.
- 7 Add 250 µl prechilled Hybridization Buffer.
- 8 Vortex briefly and then centrifuge at 280 × g for 1 minute.
- 9 Place the tube on the preheated incubator for 2 minutes.
- 10 Immediately cool on ice.
- 11 Leave on ice for 5 minutes.
- 12 [Optional] Denature and dilute a PhiX control to 1.8 pM and a 1% spike-in to the final library.
- 13 Clean the foil seal covering reservoir #16 using a low-lint tissue.
- 14 Pierce the seal with a clean 1 ml pipette tip.
- 15 Add 500 µl prepared libraries into reservoir #16.

## Perform a Sequencing Run

- 1 From the Home screen, select **Sequence**.
- 2 Enter your user name and password.
- 3 Select **Next**.
- 4 Select a run name from the list of available runs.
- 5 Select **Next**.
- 6 Open the flow cell compartment door.
- 7 Press the release button to the right of the flow cell latch.
- 8 Place the flow cell on the flow cell stage over the alignment pins.
- 9 Close the flow cell latch to secure the flow cell.
- 10 Close the flow cell compartment door.
- 11 Open the reagent compartment door.
- 12 Slide the reagent cartridge into the reagent compartment until the cartridge stops.
- 13 Remove the spent reagents bottle from the compartment.
- 14 Discard the contents and slide the empty spent reagents bottle into the compartment.
- 15 Close the compartment door and select **Next**.
- 16 Confirm run parameters.
- 17 Select **Next**.
- 18 When the automated check is complete, select **Start**.
- 19 Monitor run progress, intensities, and quality scores as metrics appear on the screen.

## View Analysis Results

- 1 From the Local Run Manager dashboard, click the run name.
- 2 From the Run Overview tab, review the sequencing run metrics.
- 3 [Optional] Click the **Copy to Clipboard**  icon for access to the output run folder.
- 4 Click the Sequencing Information tab to review run parameters and consumables information.
- 5 Click the Samples and Results tab to view the analysis report.
- 6 [Optional] Click the **Copy to Clipboard**  icon for access to the Analysis folder.