

# PMGC GENOMIC DNA EXTRACTION FOR shRNA AND sgRNA SCREENS

## **Genomic DNA (gDNA) Extraction**

gDNA extraction is performed using the QIAamp Blood Maxi kit (Cat#51194) essentially as described in the kit manual. Follow the instructions below, using the indicated volumes and refer to the kit manual for extra details if required.

**This protocol is optimized for cell pellets containing 20-50x10<sup>6</sup> cells.**

\*\*\*It is very important to use a swinging-bucket rotor in a centrifuge that is capable of attaining the required g-force. Failure to do so will result in low yield and dirty genomic DNA\*\*\*

Prepare 70°C water bath with rack for your tubes.

Ensure that buffers are prepared and ready to use.

Thaw cell pellets in 37°C water bath for 5-10 min.

Label samples and pre-label tubes and columns to be used in extraction.

**Use plugged pipette tips for all steps** – contamination of the gDNA is to be avoided!

- Add 4.5ml of sterile PBS to each cell pellet. Seal tubes tightly and vortex thoroughly to disperse cells. Pipette to disrupt cell clumps if needed – no clumps of cells can remain or the prep will not work well.
- Add 500µl of Qiagen protease solution to each sample. Mix briefly by swirling.
- Add 6ml Buffer AL. Cap tubes and mix by inversion for 2 min.
- Incubate tubes in 70°C water bath for 15 min.
- Let tubes cool to ~40°C, add 5ml 100% ethanol. Mix by shaking and inverting for 2 min.
- Carefully pour all of the solution into a pre-labeled QIAamp maxi column placed in the provided 50ml centrifuge tubes. Take care not to spill onto the rim of the column.
- Centrifuge tubes at 1850g for 3 min.
- Remove the QIAamp maxi column, vacuum aspirate the filtrate (**do not** pour out), and place the column back into the tube.
- Add 5ml Buffer AW1 to the column, being careful to not spill onto the rim.
- Cap the tubes and centrifuge at 4500g for 2 min.
- Add 5ml Buffer AW2 to the column, being careful to not spill onto the rim.
- Cap the tubes and centrifuge at 4500g for 20 min.

- Check the tubes – if any buffer remains on the inside edge of the column, or if the filter appears to be wet, uncap the columns and dry in a warm incubator for ~15 minutes (or remove by pipette/aspiration – for wet edge only, NOT wet filter).
- Place the QIAamp maxi columns into clean 50ml centrifuge tubes, and discard the filtrate and the previous tubes.
- Add 600µl of room temperature Buffer AE directly onto the membrane and cap the tubes.
- Incubate the tubes at room temperature for 10 min, then centrifuge at 4500g for 3 min.
- Pipette the flow-through (less than 600µl) back onto the membrane, and dispense an additional 150µl of Buffer AE onto the membrane.
- Incubate the tubes at room temperature for 10 min, then centrifuge at 4500g for 10 min.
- Collect flow through and transfer 500µl to eppendorf tubes (save extra eluate in case you need it).
- Quantitate DNA and measure purity by spectrophotometry (we suggest using a Nanodrop instrument. Use 4µl of 2mg/ml DNase-free RNase A for every ~30µg, mix well by pipetting. Incubate at 37°C for 1hr.

## Genomic DNA Precipitation

- Estimate total quantity of gDNA being precipitated based on volume and measured concentration. Assume ~75% of the concentration that you got from the nanodrop reading prior to RNase A treatment was from DNA.
- Add 5M NaCl to a final concentration of 0.2M and 2 volumes of -20°C 96-100% ethanol. Do **NOT** use sodium acetate as residual acetate interferes with our PCR reaction.
- Invert tubes 15 times, then spin at 13,000rpm for 15 minutes at 4°C in a table-top centrifuge.
- Aspirate supernatant, being careful to avoid DNA pellet.
- Add 500µl -20°C 70% ethanol.
- Wash pellet by inverting tube 10 times, then centrifuge at 13,000rpm for 10 minutes at 4°C.
- Aspirate supernatant. Pulse-spin down any remaining liquid and aspirate it, being careful to avoid the DNA pellet.
- Air dry pellet ~ 5-10 minutes (not longer unless it is still wet) and resuspend with repeated pipetting in buffer EB (10mM Tris-HCl pH 7.5) to a final estimated concentration of 450ng/µl.
- Heat samples at 50°C for 1 hour. Pipette and vortex repeatedly to fully resuspend/solubilize the DNA. **Make sure that DNA is fully solubilized with no lumps or clumps.**

### IMPORTANT:

- Check concentration and quality of DNA samples via Nanodrop and dilute to 400ng/µl.
- OD 260/230 ratio **SHOULD** be >2.0 and OD 260/280 ratio **SHOULD** be 1.8 +/- 0.05.

## Sample Drop-off / Shipping

If dropping off samples: Please **schedule your drop off date and time in advance** with your PMGC contact person.

- Your PMGC contact will meet you at the **9<sup>th</sup> floor elevator lobby** of the Princess Margaret Cancer Research Tower (PMCRT) at your pre-arranged time. PMCRT is the East Tower of the MaRS building, near the corner of College and Elizabeth Street entrance.
- Email or call/text when you are at the designated meeting area and your PMGC contact will come to collect the samples.
- REMINDER: Transport samples using appropriate means of storage (e.g. on dry ice for frozen samples, wet ice for fresh samples). Please confirm with PMGC if any questions.

If shipping samples: Please ship out on **Monday/Tuesday** to prevent weekend delays. Place a generous supply of dry ice to ensure dry ice will remain for the duration of the delivery time. For international clients, we recommend shipping with [World Courier](#). Within Canada, or if shipping DNA/RNA, we recommend FedEx Next Day Priority services.

Shipping address:

Attn: (insert PMGC contact person)  
Princess Margaret Genomics Centre  
101 College St.  
PMCRT, Rm 9-601A  
Toronto, Ontario M5G 1L7  
Canada