


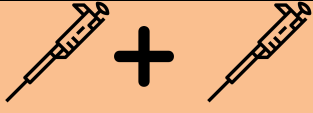
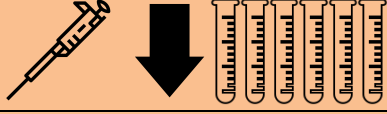

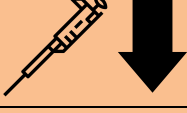


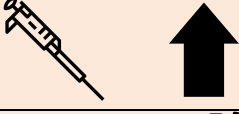
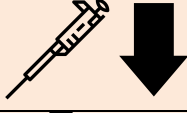
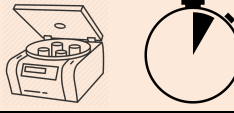
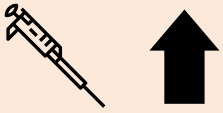











Whole blood processing kit

Graphical protocol

Staining of fixation sensitive epitopes

1. <input type="checkbox"/>		Prepare staining cocktail containing antibodies targeting sensitive epitopes
2. <input type="checkbox"/>		Mix blood with the Ab-cocktail
3. <input type="checkbox"/>		Incubate at 5-8°C/RT for 30-60 min
		<p>Calculate required volume and prepare 1x Fixation Buffer 5-10 min before the end of the staining step:</p> <ul style="list-style-type: none"> ↓ 1 part of 2x Fix-Concentrate ↓ 1 part of Fix-Diluent <p>Equilibrate at room temperature for 5 min.</p>
4. <input type="checkbox"/>		Dispense fixation buffer to tubes
5. <input type="checkbox"/>		Incubate sample at +20°C for 15 minutes. Vortex multiple times throughout.
6. <input type="checkbox"/>	LYSIS 	Add LYSIS buffer and vortex.
7. <input type="checkbox"/>		Incubate sample at +20°C for 15-20 minutes (until the color of the solution is crystal clear red). Do not proceed without complete RBCs lysis!
8. <input type="checkbox"/>		Centrifuge sample at: 300g for 5 minutes.
9. <input type="checkbox"/>		Aspirate supernatant but keep some volume at the bottom. Do not decant!
10. <input type="checkbox"/>	WASH 	Add WASH buffer.
11. <input type="checkbox"/>		Centrifuge sample at: 300g for 5 minutes.
12. <input type="checkbox"/>		Aspirate supernatant but keep some volume at the bottom. Do not decant!

13. <input type="checkbox"/>	 #10, #11 & #12	If the pellets contain remaining RBCs (pinkish- red), repeat washing procedure (WASH buffer/ centrifugation/supernatant aspiration).	
14. <input type="checkbox"/>	<input type="checkbox"/> 14.a.	<input type="checkbox"/> 14.b.	
		 $\leq -80^{\circ}\text{C}$	
	Proceed to staining of remaining epitopes/ICS.	Store samples in CRYO#20 and freeze at $\leq -80^{\circ}\text{C}$.	
15. <input type="checkbox"/>		   37°C	
		Thaw cryogenically preserved sample using 37°C water bath (≈ 1 min)	
16. <input type="checkbox"/>	<input type="checkbox"/> 16.a.	<input type="checkbox"/> 16.b.	
	<u>Mass cytometry</u> Wash once or twice with staining buffer	<u>Flow cytometry</u> Acquire sample without washing or stain directly in CRYO#20 (17.b.).	
17. <input type="checkbox"/>	<input type="checkbox"/> 17.a.	<input type="checkbox"/> 17.b.	
	 Follow your standard mass cytometry staining protocol	 Adjust volume by short spin 400g/2-3 min and aspirate supernatant. Leave 50-100 μl Proceed to staining of remaining epitopes/ICS.	
18. <input type="checkbox"/>	ACQUIRE		

Examples of recommended processing conditions						
Blood sample volume	Fix buffer volume	LYSIS buffer volume	WASH buffer volume	Recommended processing tube size		CRYO#20 volume
				First wash	All consequent washes	
100 μl	1.0 ml	4.0 ml	4.0 ml	10 – 15 ml	5-15 ml	100 μl
250 μl	2.5 ml	10 ml	10 ml	15 ml	15 ml	100 μl
500 μl	5.0 ml	20.0 ml	20.0 ml †	50 ml *	15 ml	250 μl
1.0 ml	10.0 ml	39.0 ml	40.0 ml †	50 ml *	15 ml	500 μl

* To achieve optimal cell yields execute only first spin after Fix&Lyse step in 50 ml tube, aspirate to mark 5 ml or above (not below 5 ml mark), transfer to 15 ml tube and continue with altered WASH buffer volumes.

† In case you decide to run all consequent processing steps in 15 ml tube, use 14 ml of WASH buffer and adjust number of washing steps based on pellet color. Typically, only one more washing step is required.