

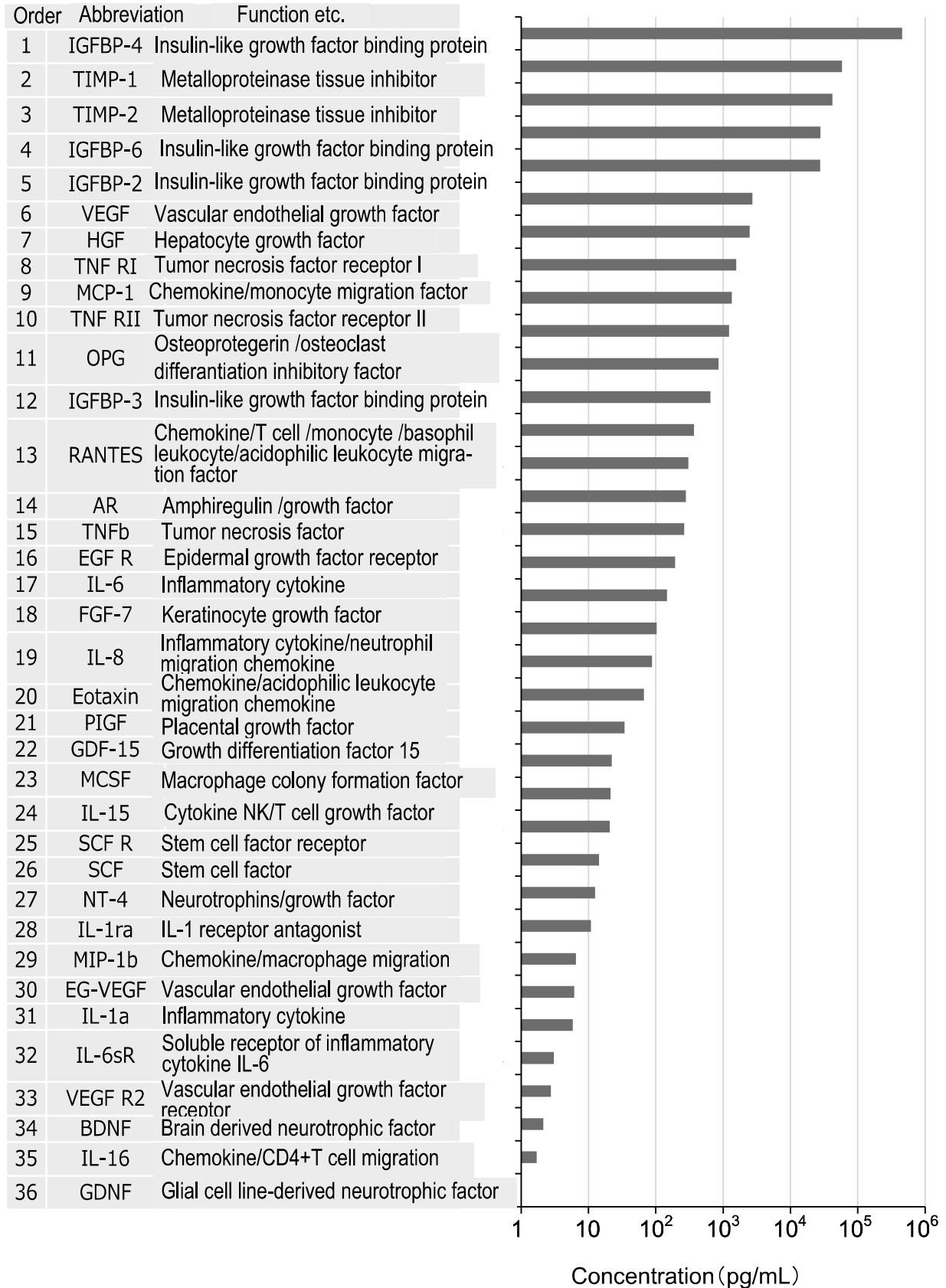
Fig. 1

Fig. 2

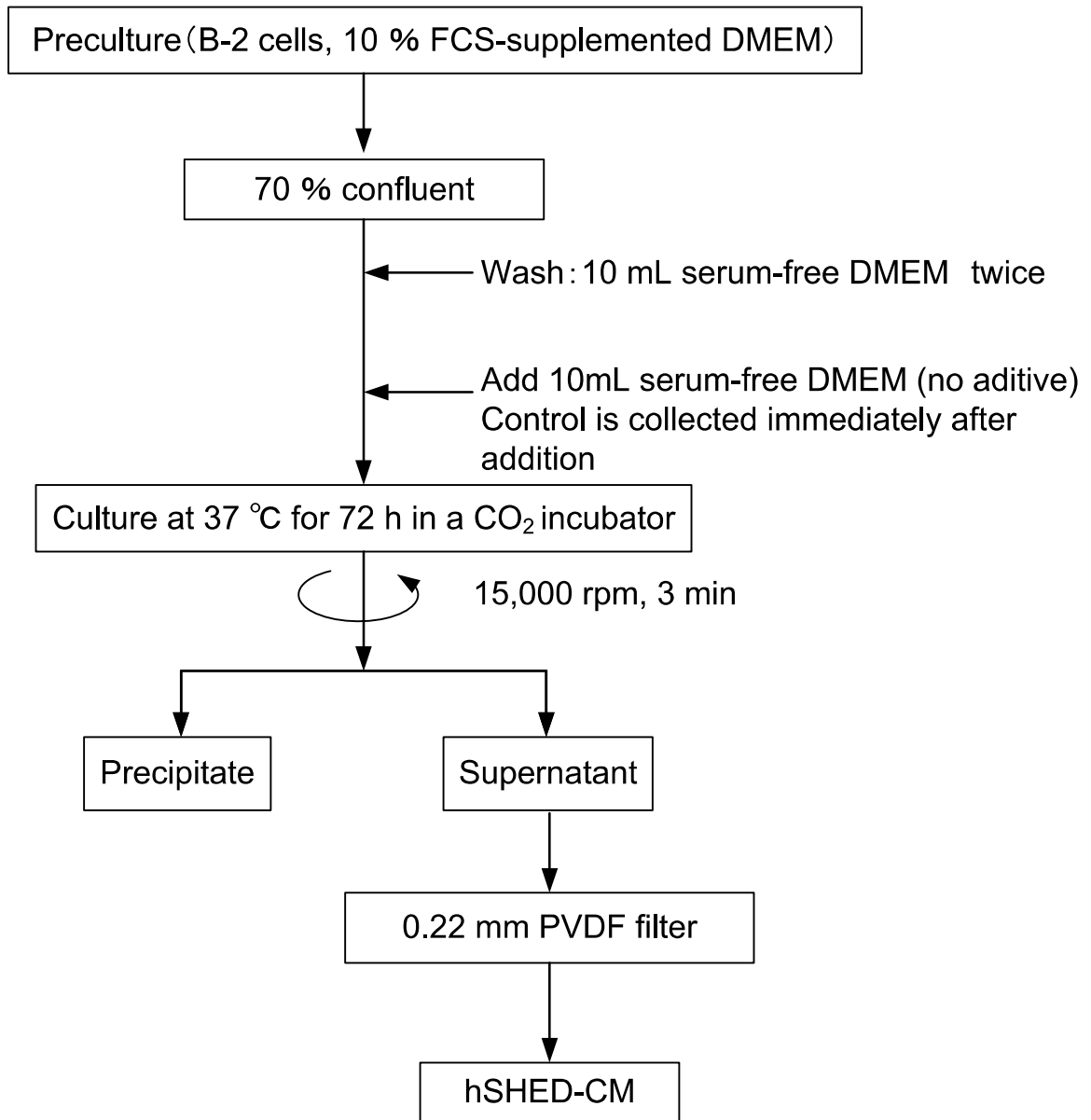


Fig. 3

<Quantitative expression analysis>

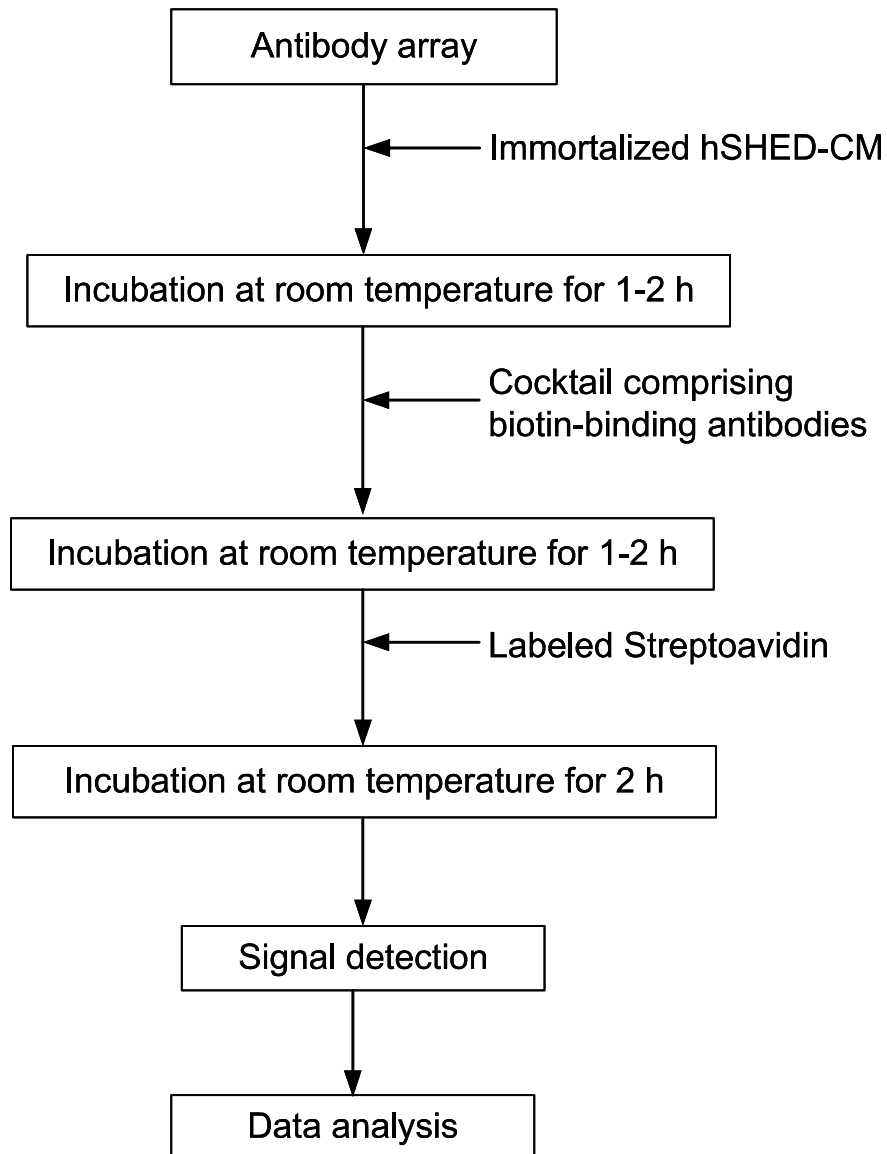


Fig. 4

<Qualitative expression analysis>

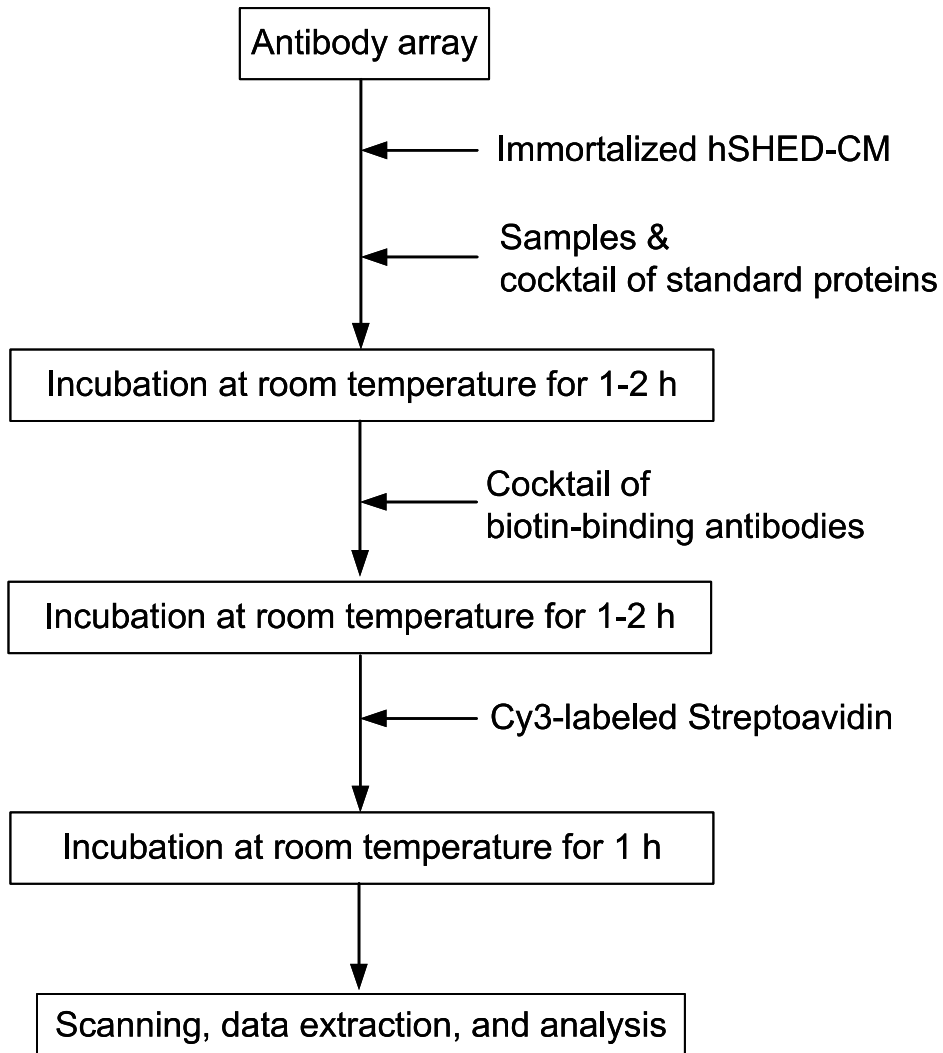


Fig. 5

<Protocol for microRNA array analysis>

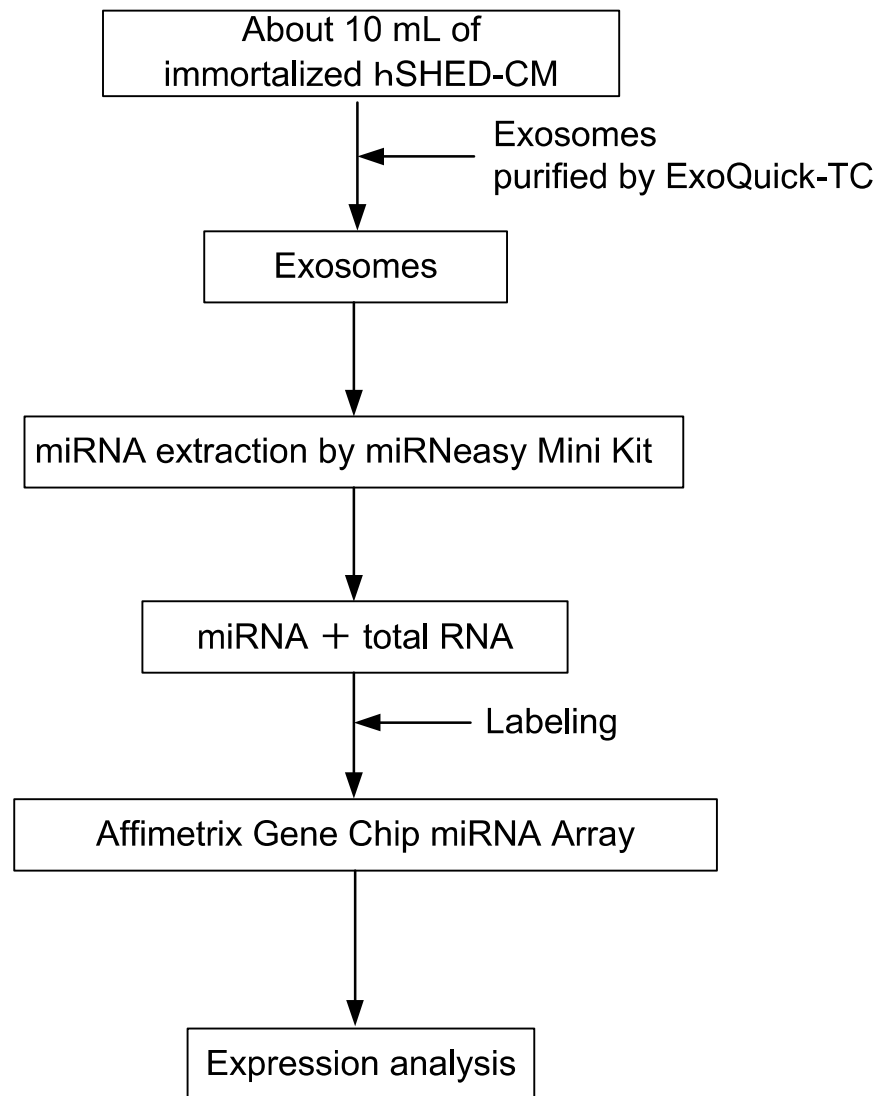


Fig. 6

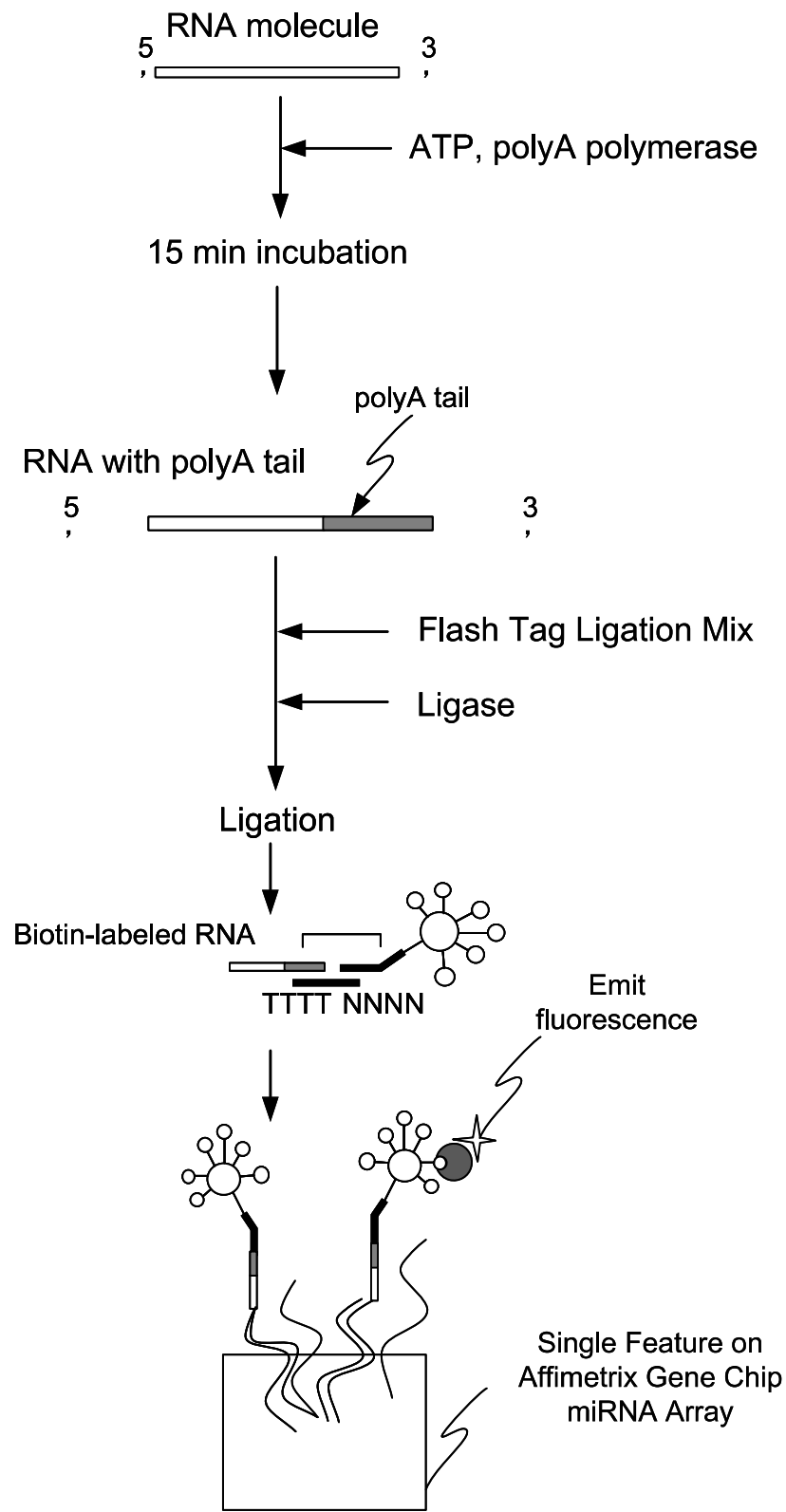


Fig. 7

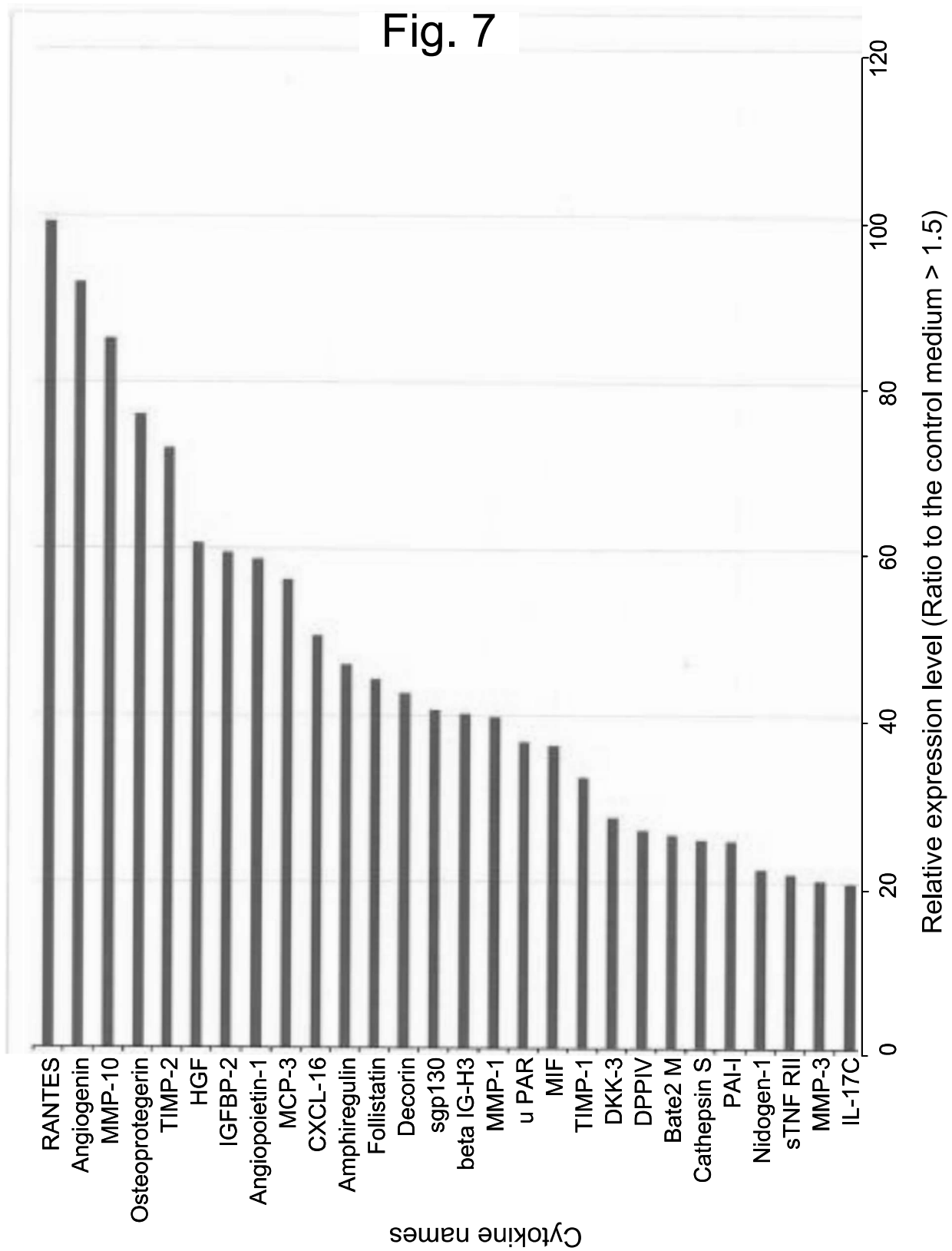


Fig. 8

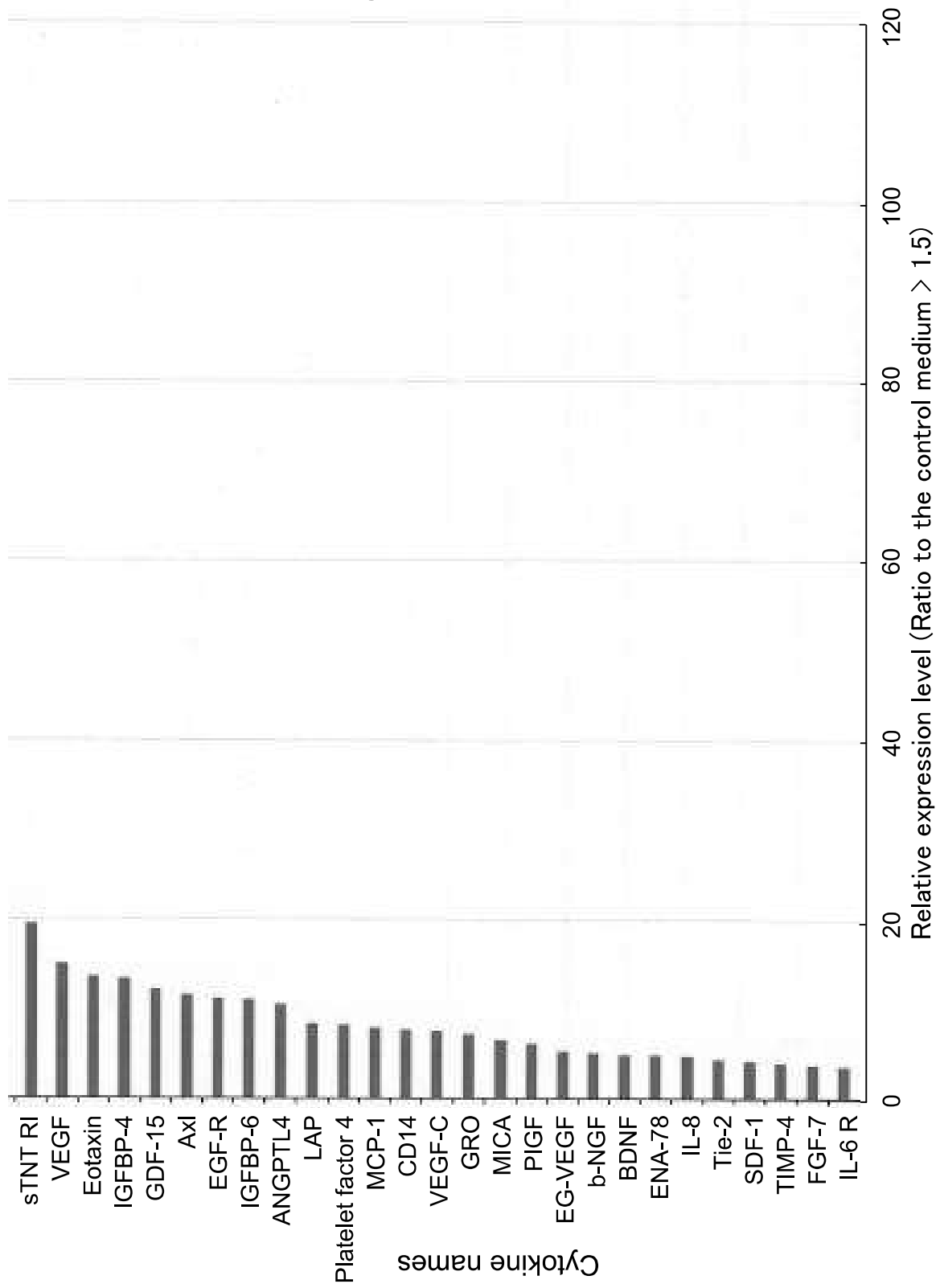


Fig. 9

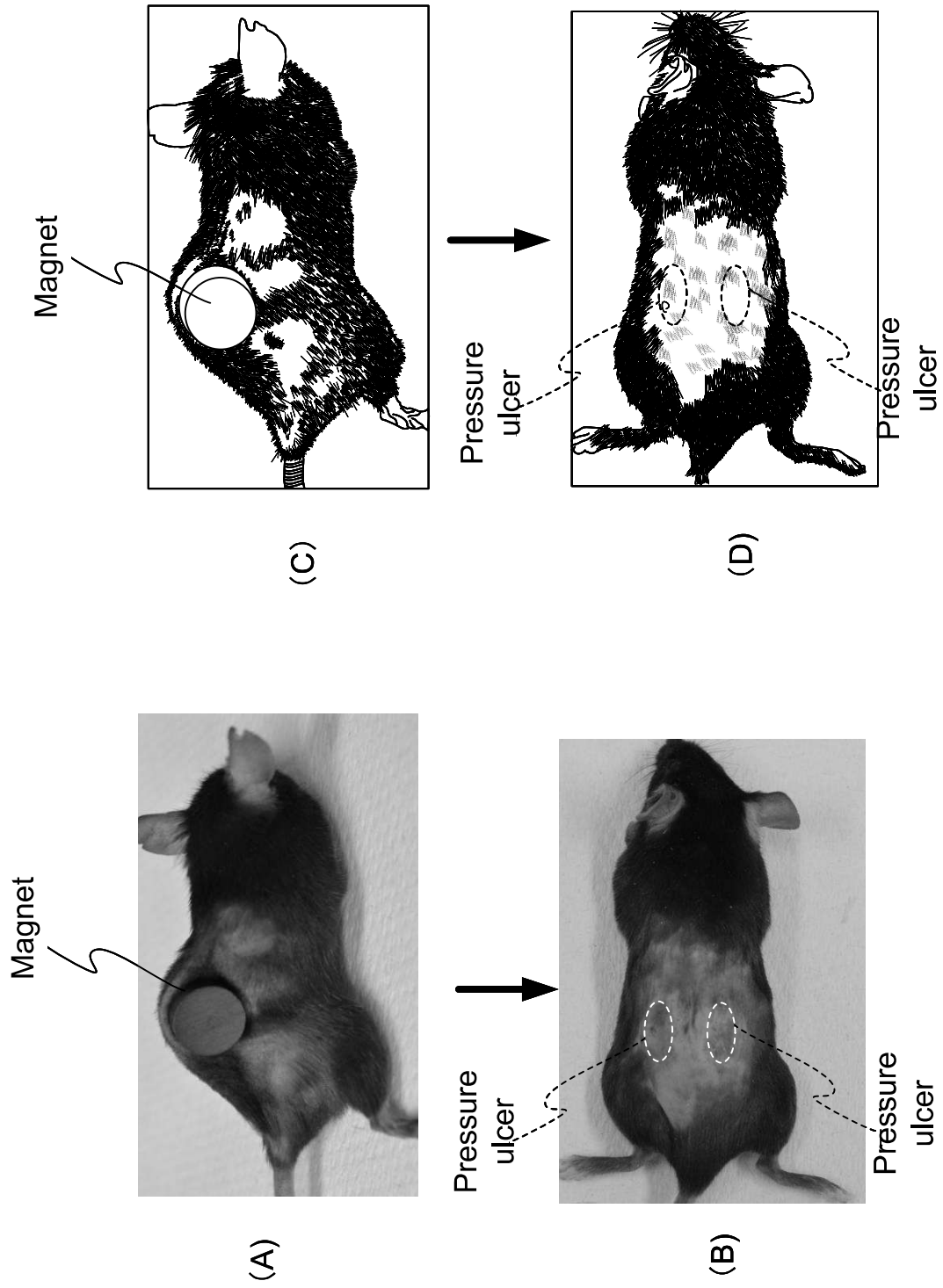


Fig. 10

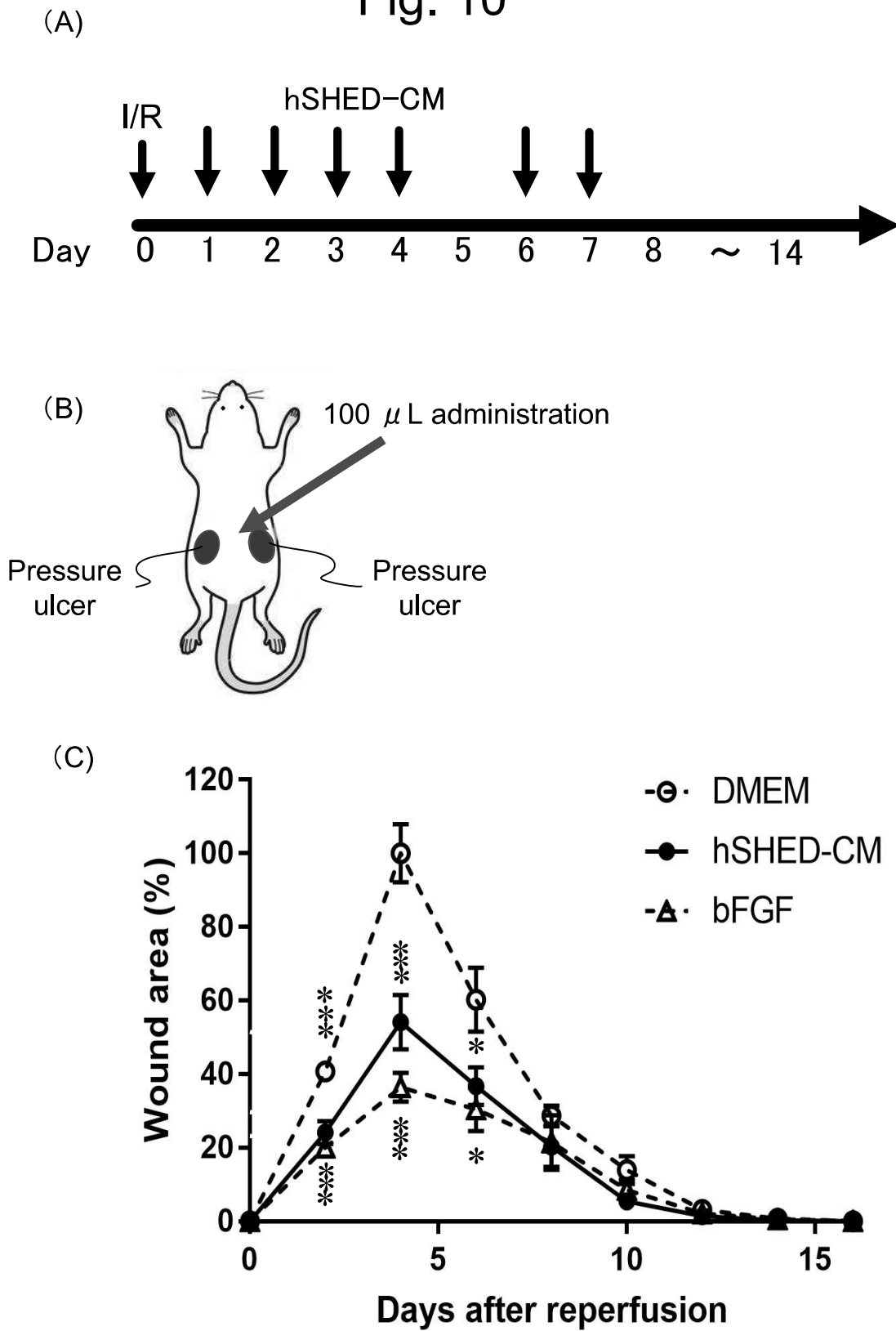


Fig. 11

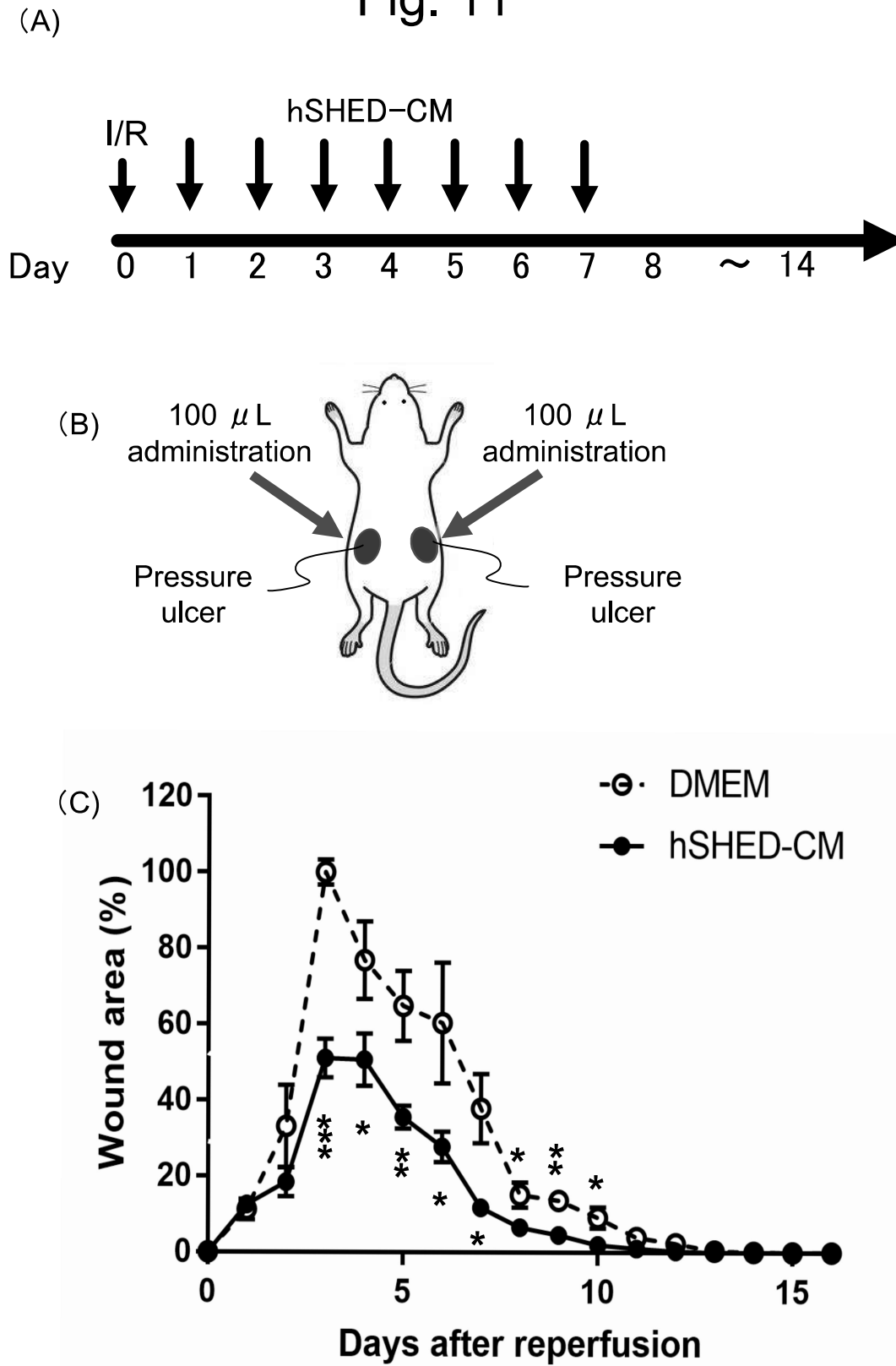


Fig. 12

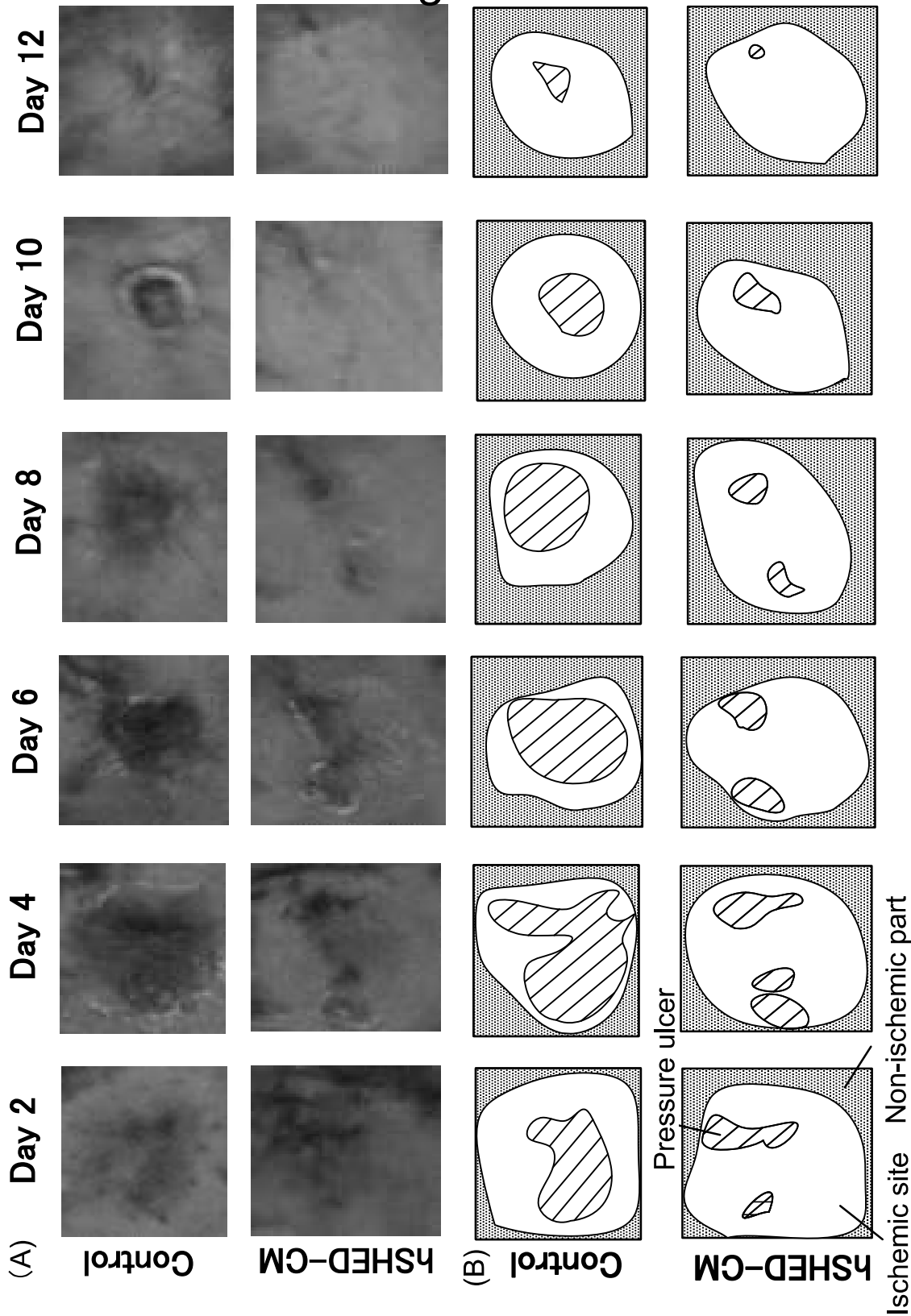
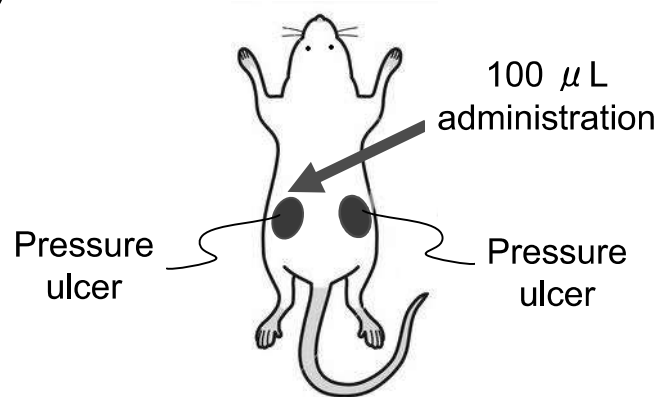
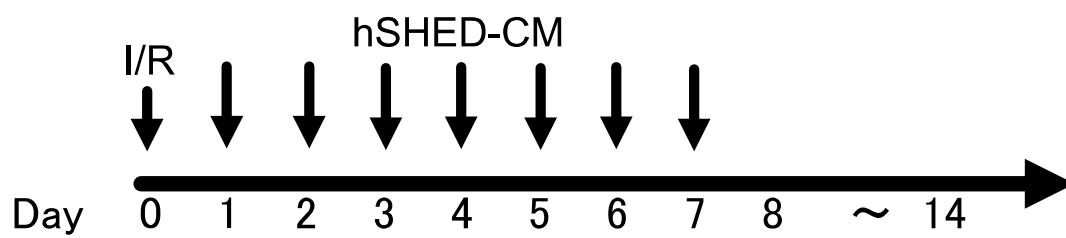


Fig. 13

(A)



(B)



(C)

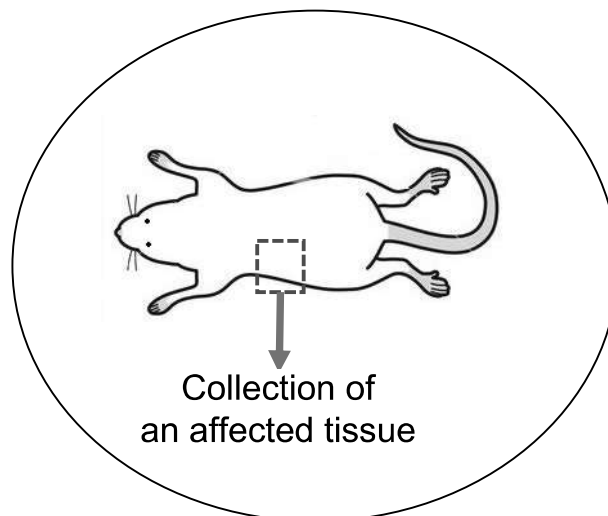
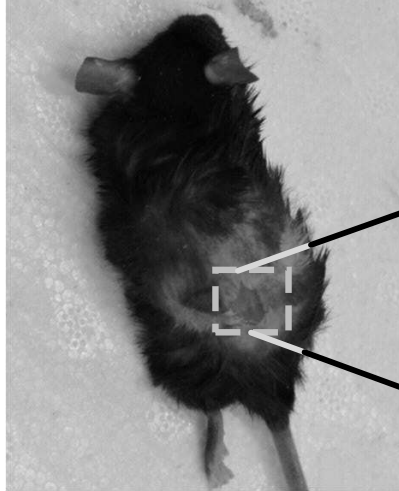


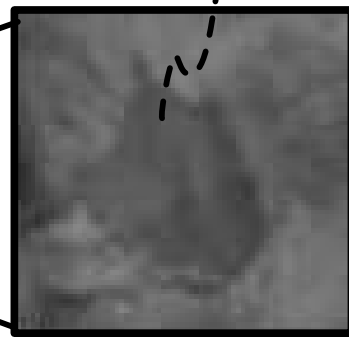
Fig. 14

(A) Condition after the affected tissue collection

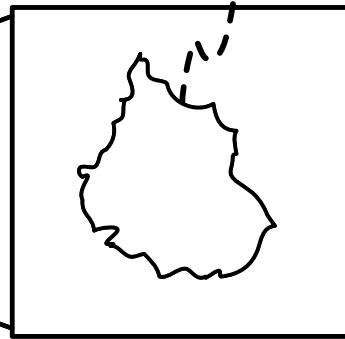
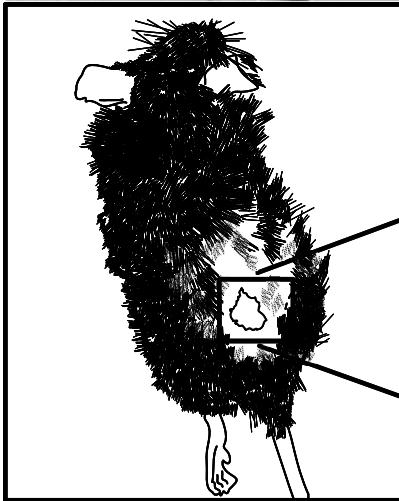


(B)

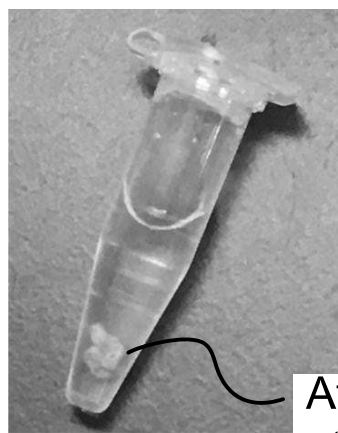
Collected site of the affected tissue



Collected site of the affected tissue

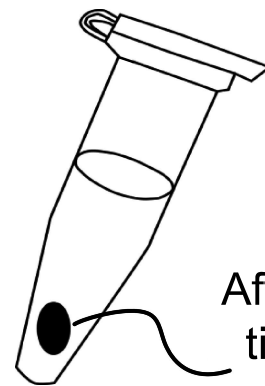


(C)



Formalin

Affected tissue



Affected tissue

Fig. 15

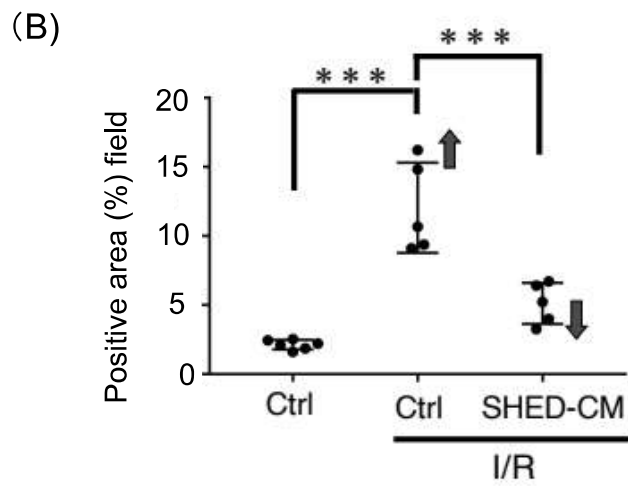
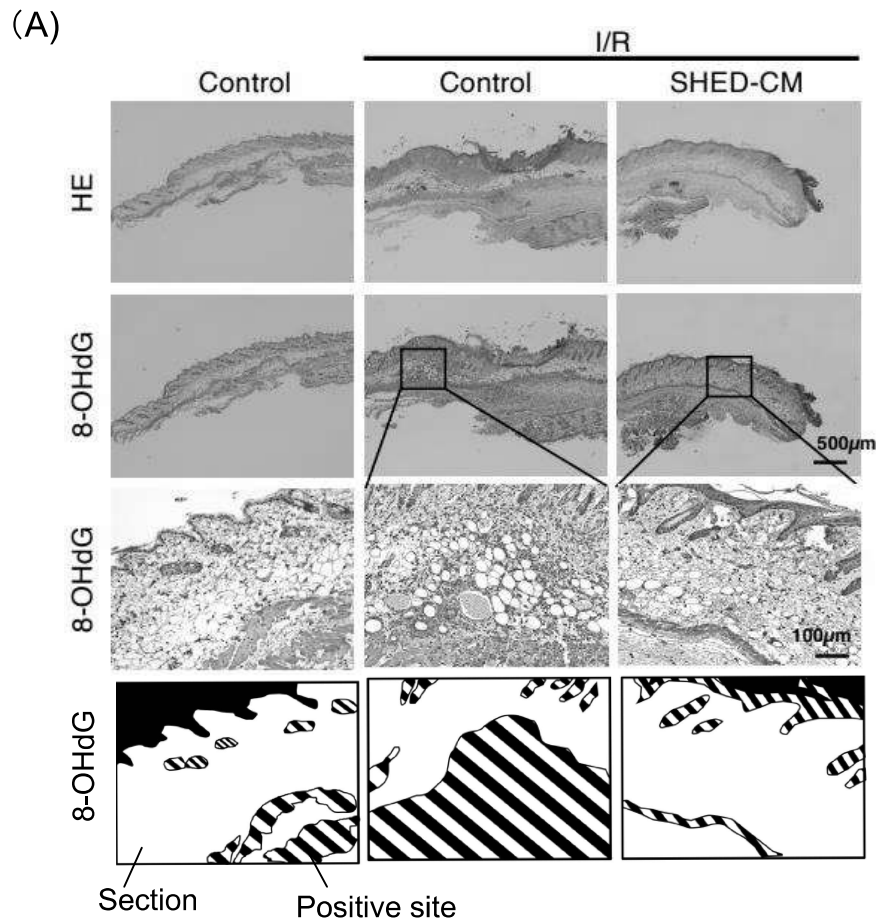
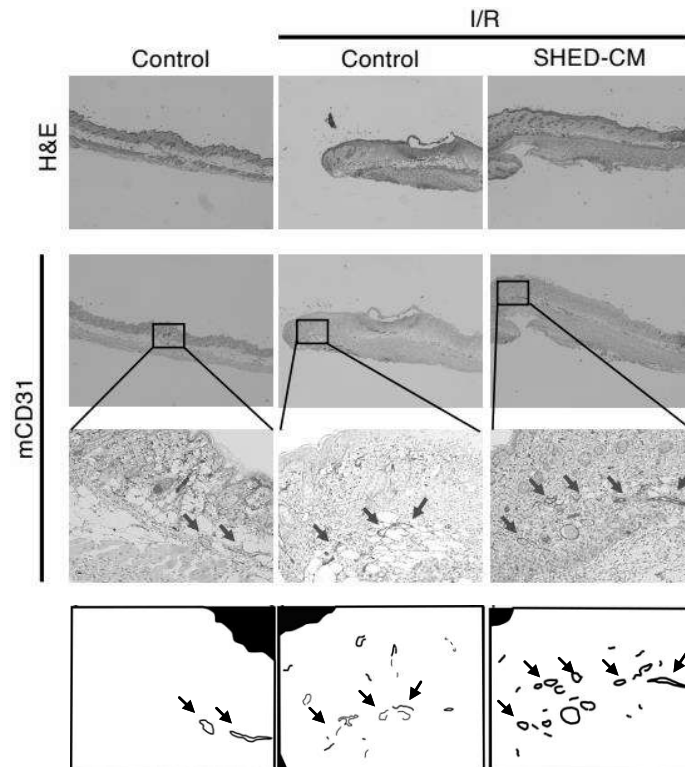


Fig. 16

(A)



(B)

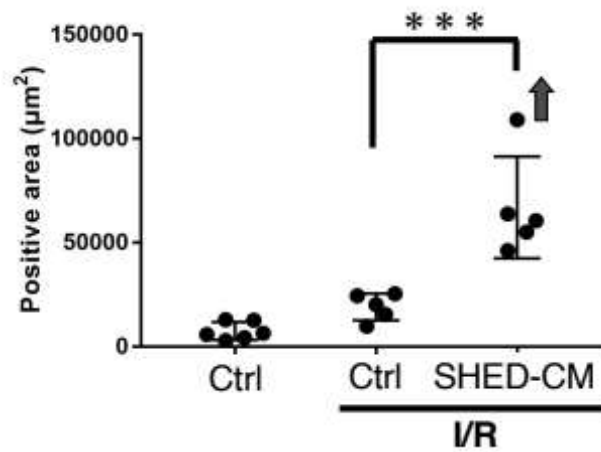
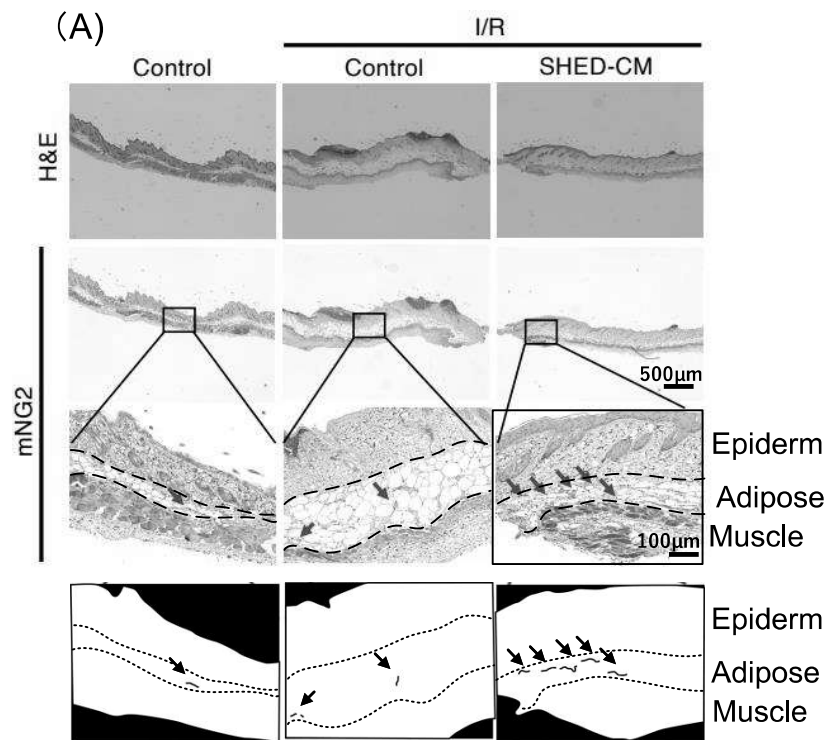
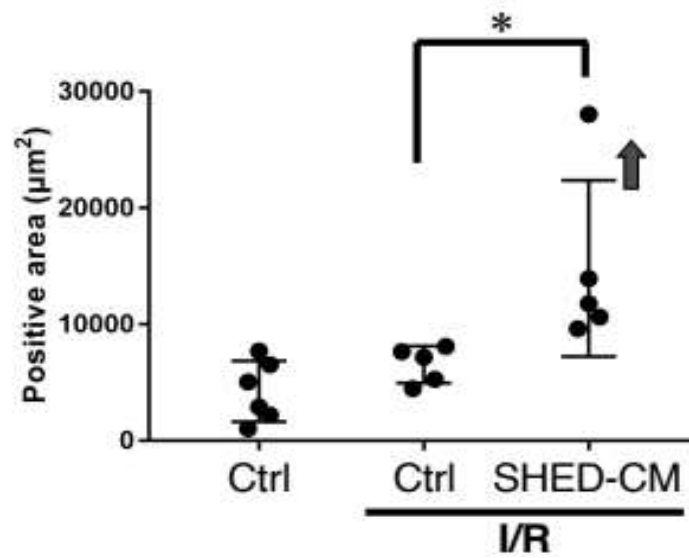


Fig. 17



(B)



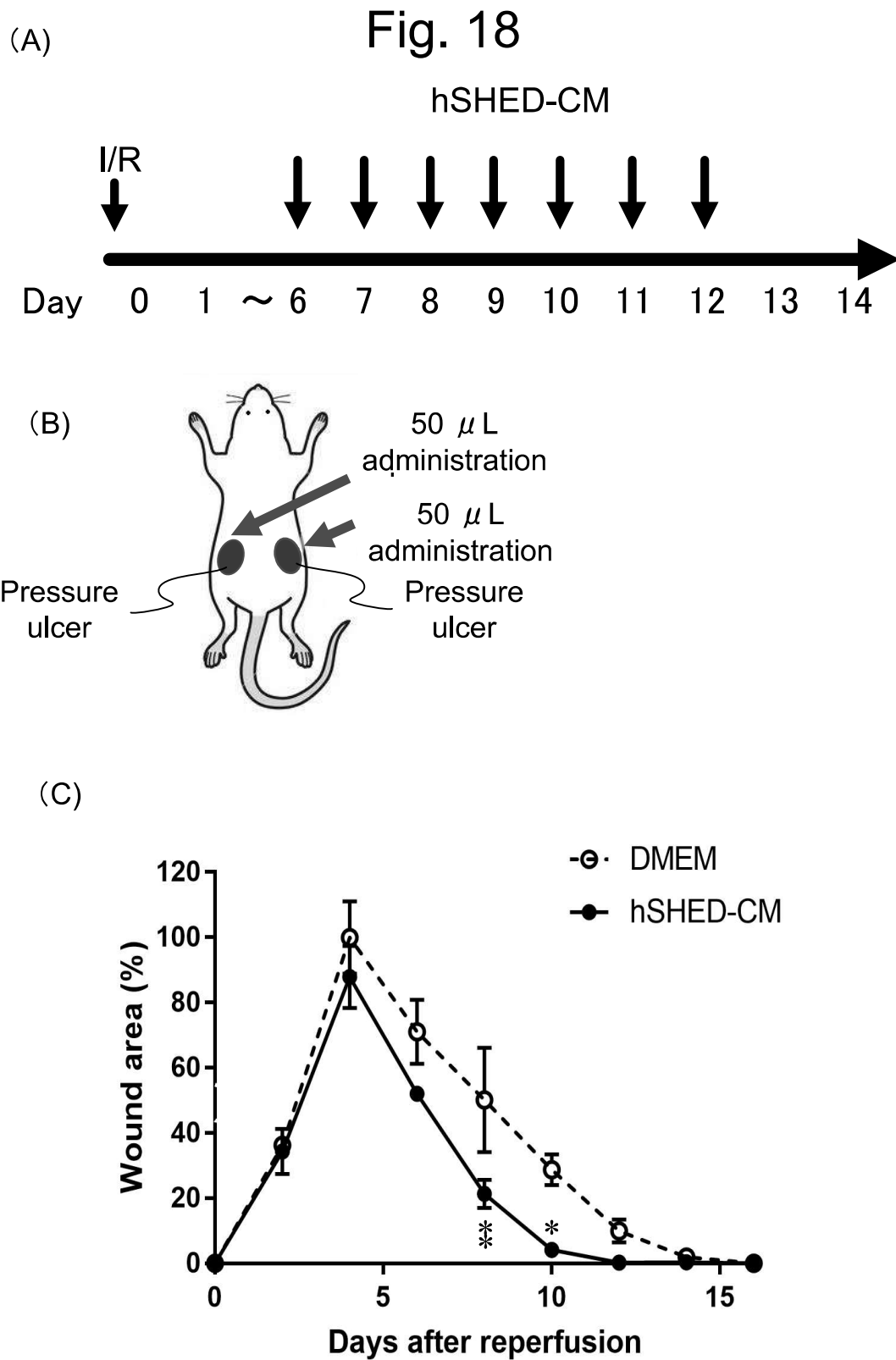


Fig. 19

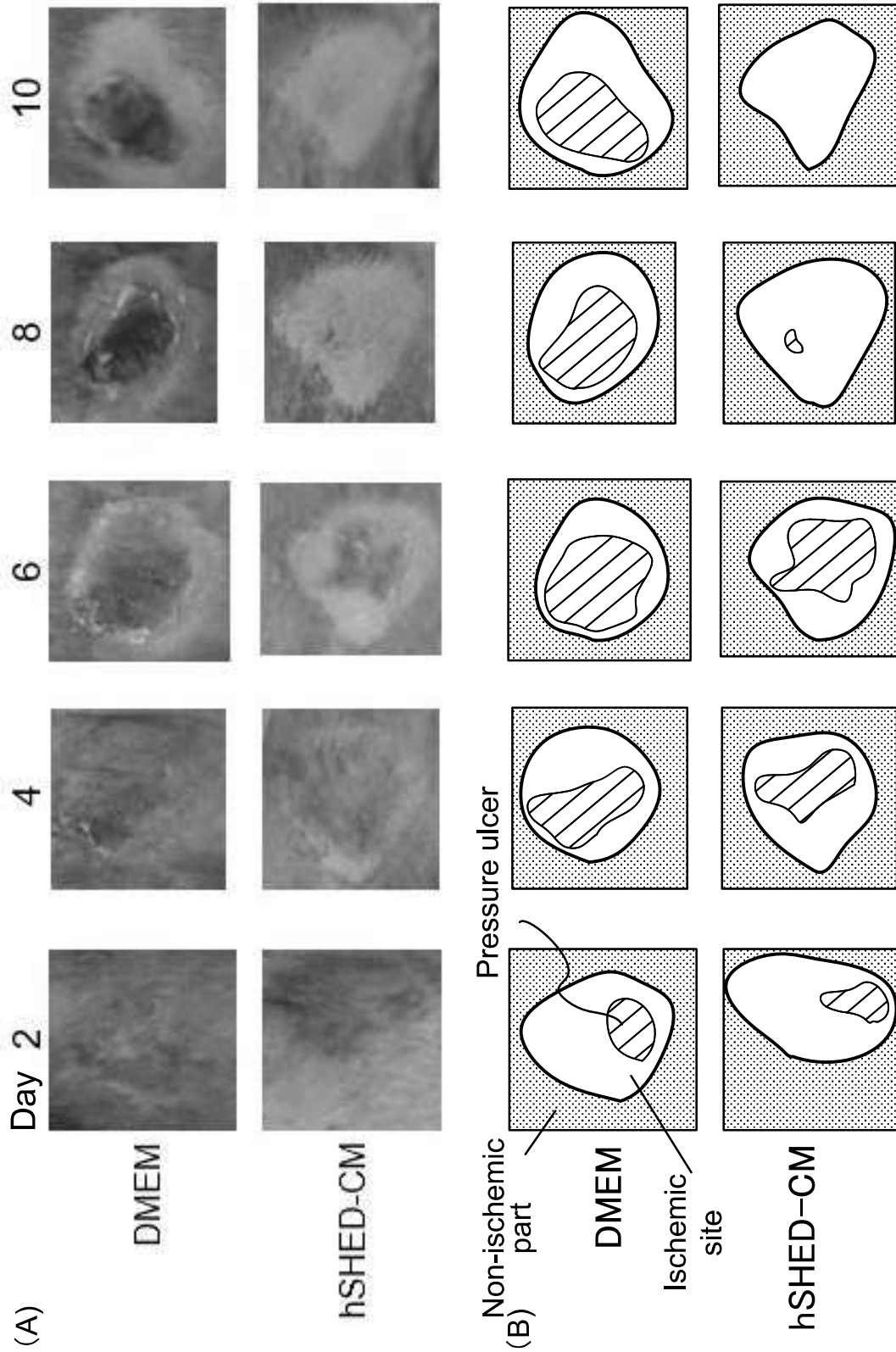


Fig. 20

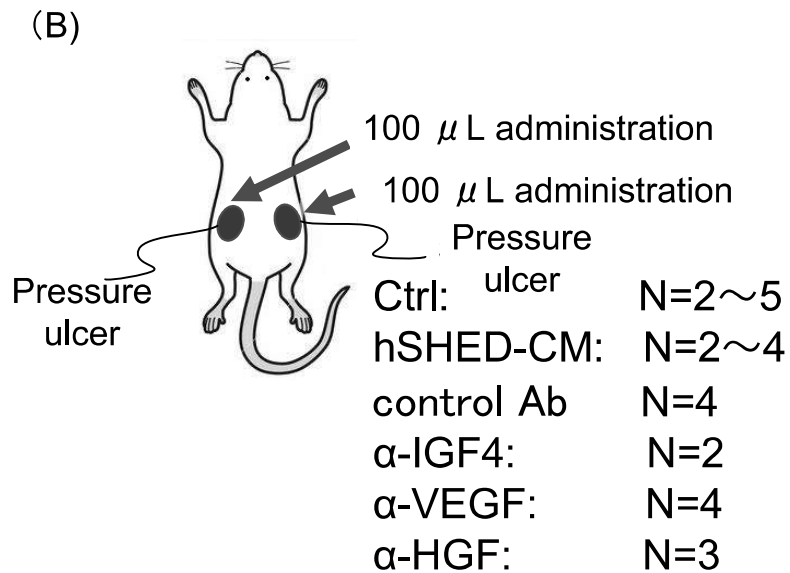
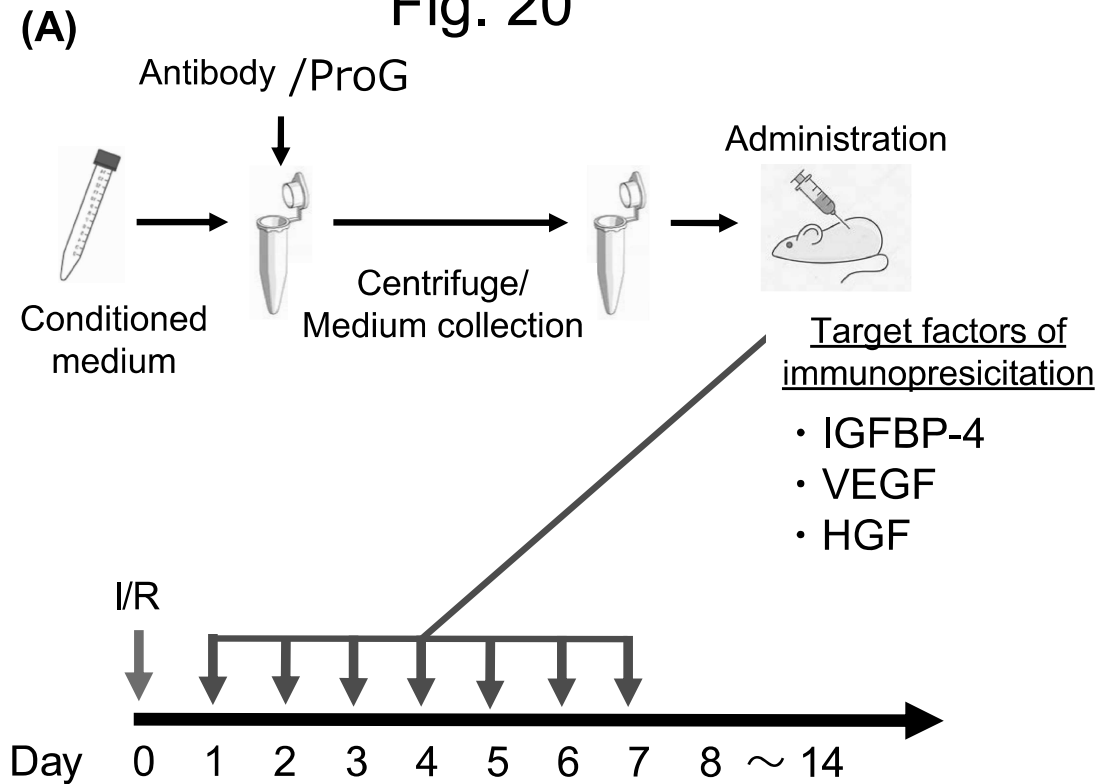
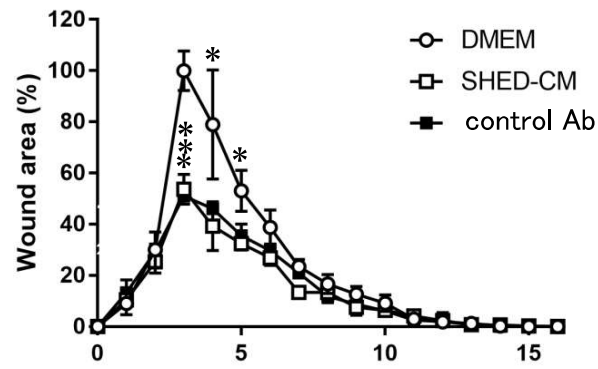
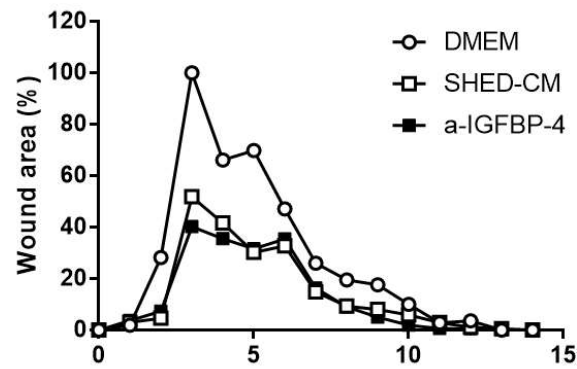


Fig. 21

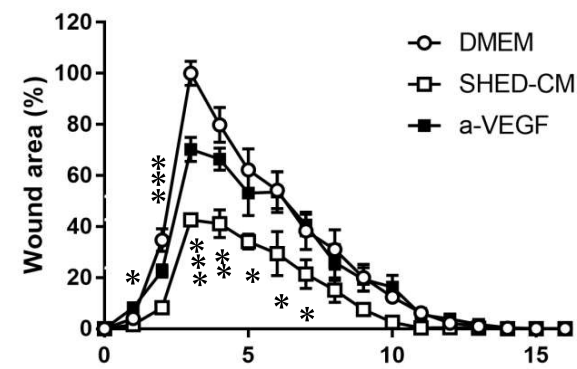
(A)



(B)



(C)



(D)

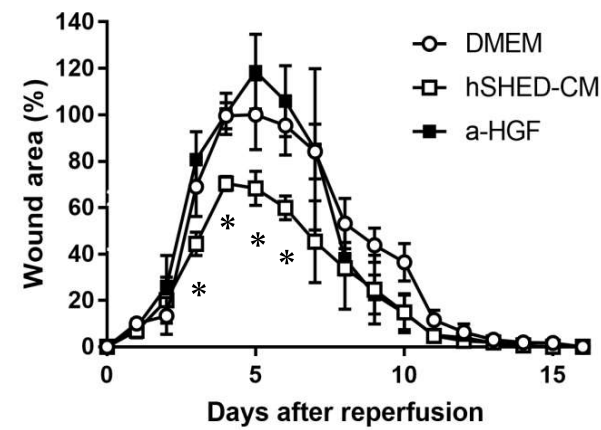
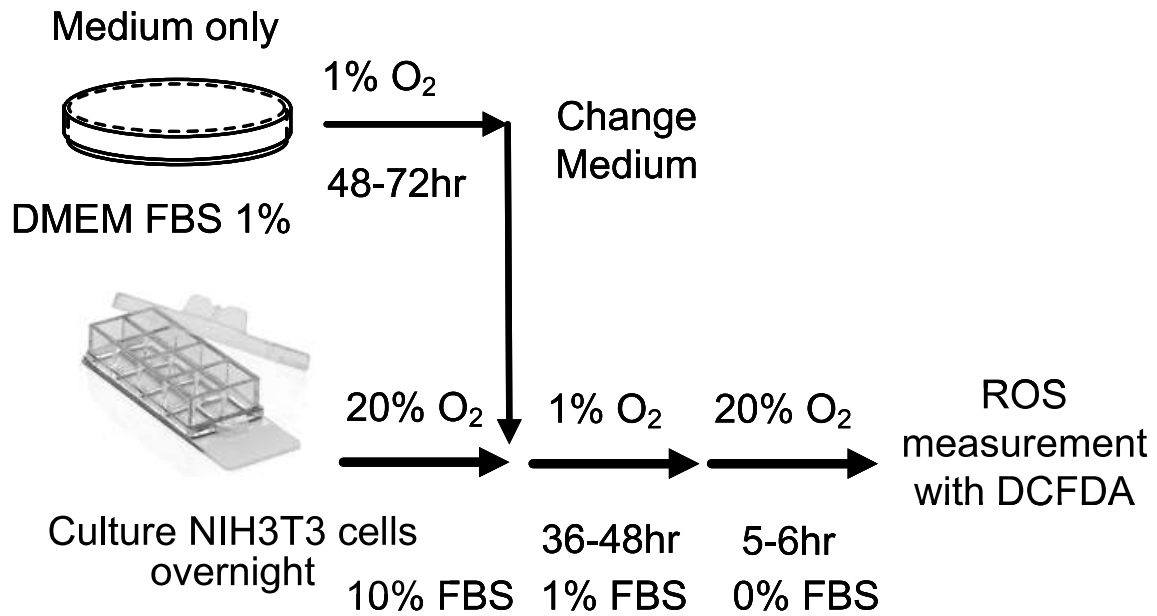


Fig. 22

Induction of ROS production
by low oxygenized-reoxygenized conditioning in NIH3T3



(A)

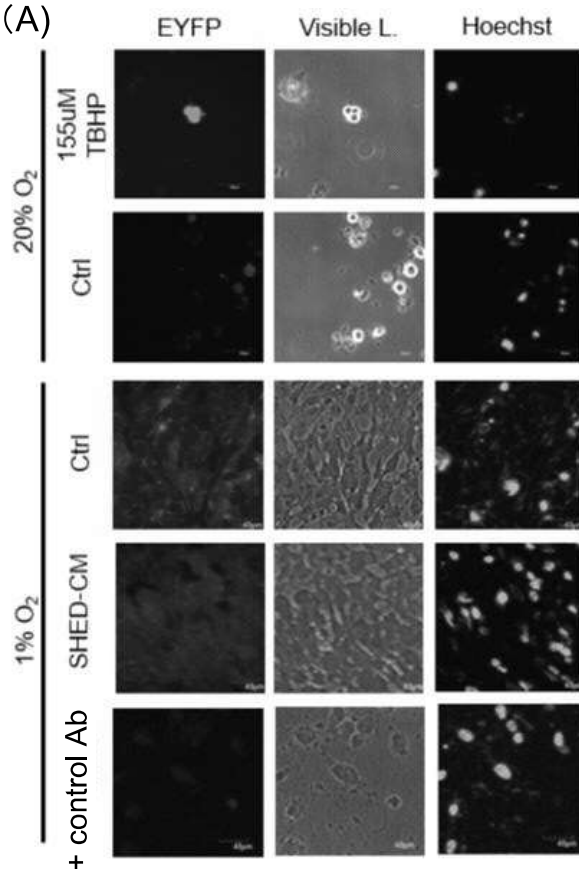
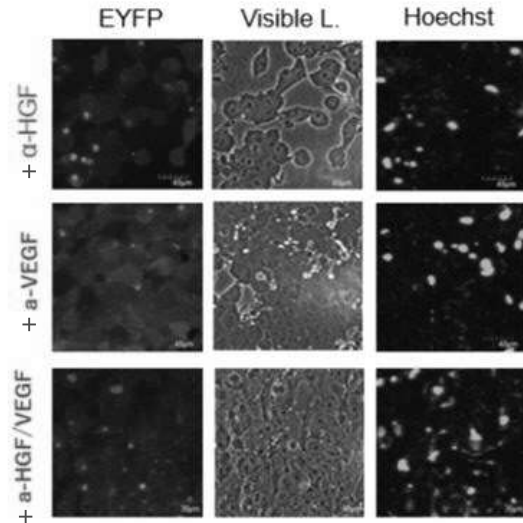
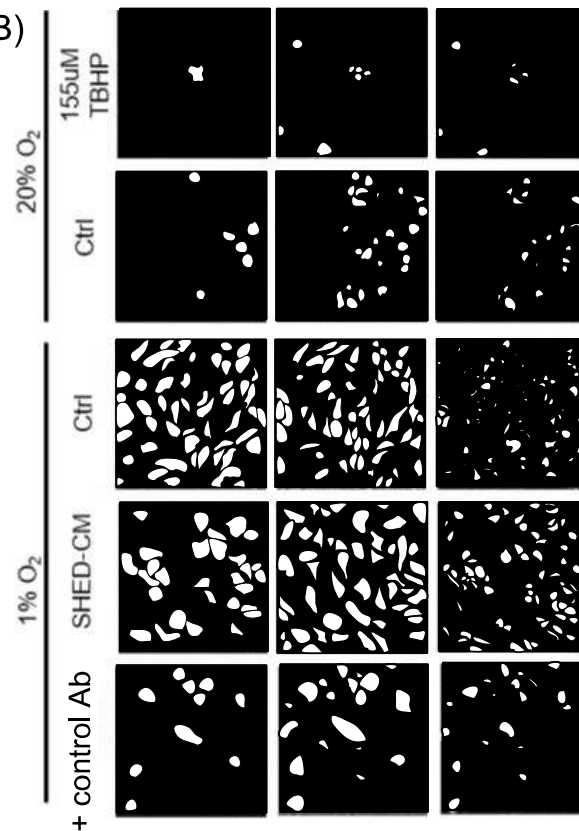


Fig. 23

Recommends: 485/535 nm
 EYFP: 480/527 nm
 FITC: 495/519 nm



(B)



Recommends: 485/535 nm
 EYFP: 480/527 nm
 FITC: 495/519 nm

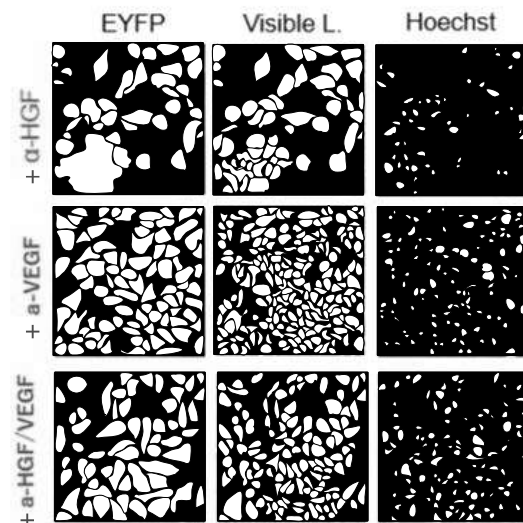


Fig. 24

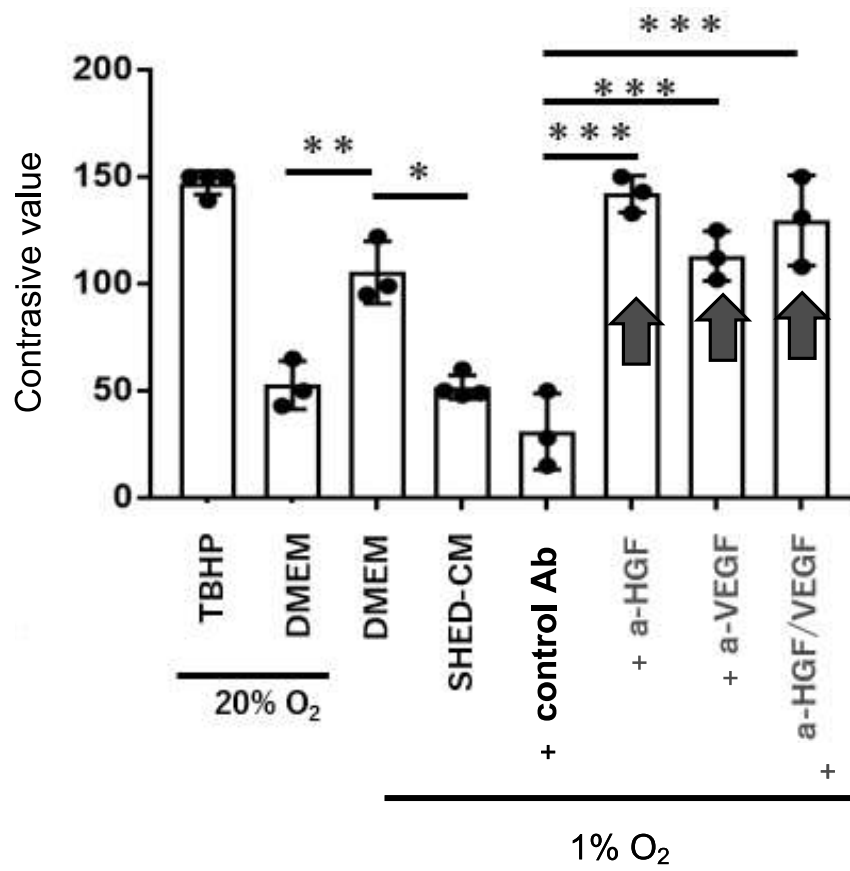


Fig. 25

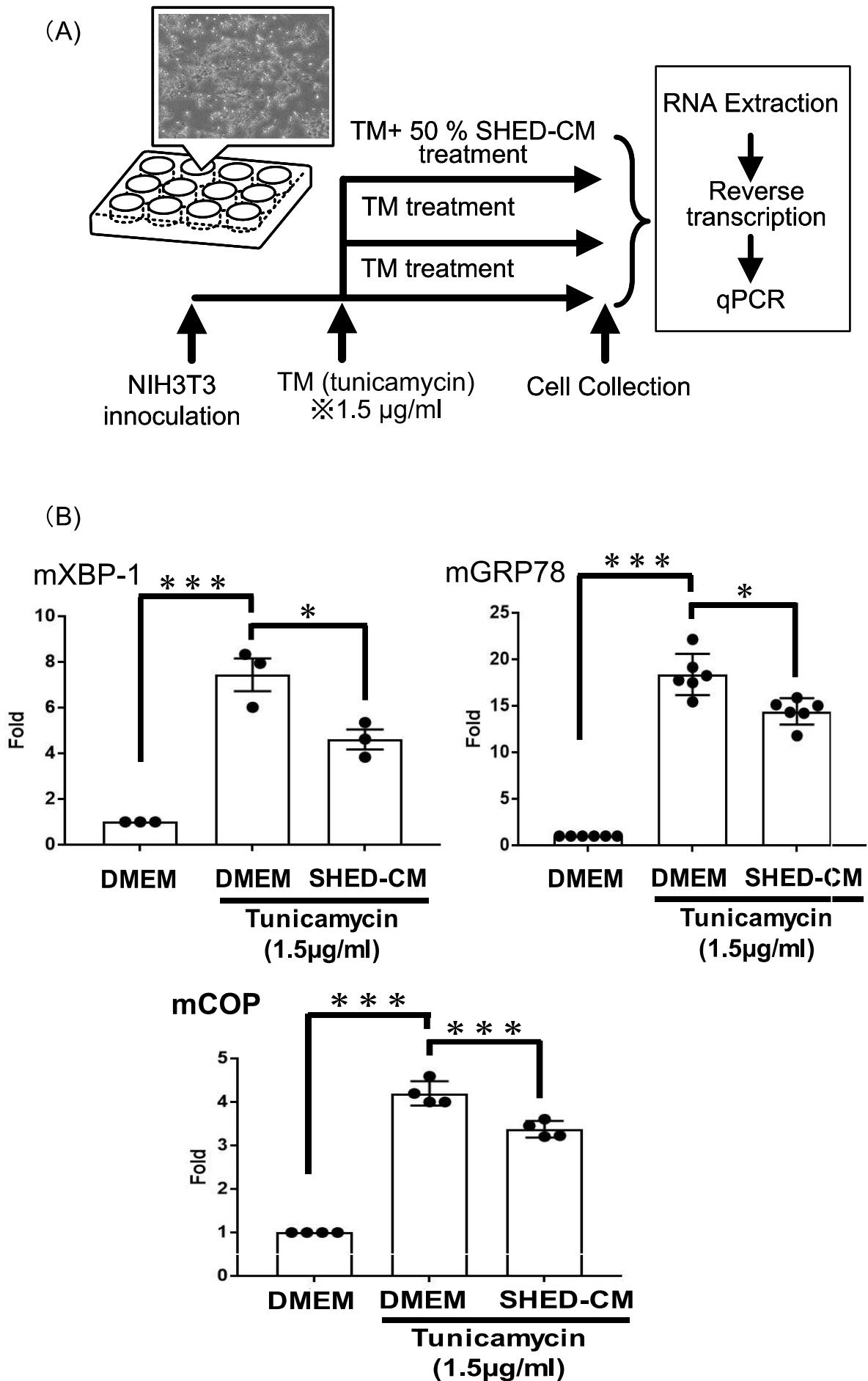


Fig. 26

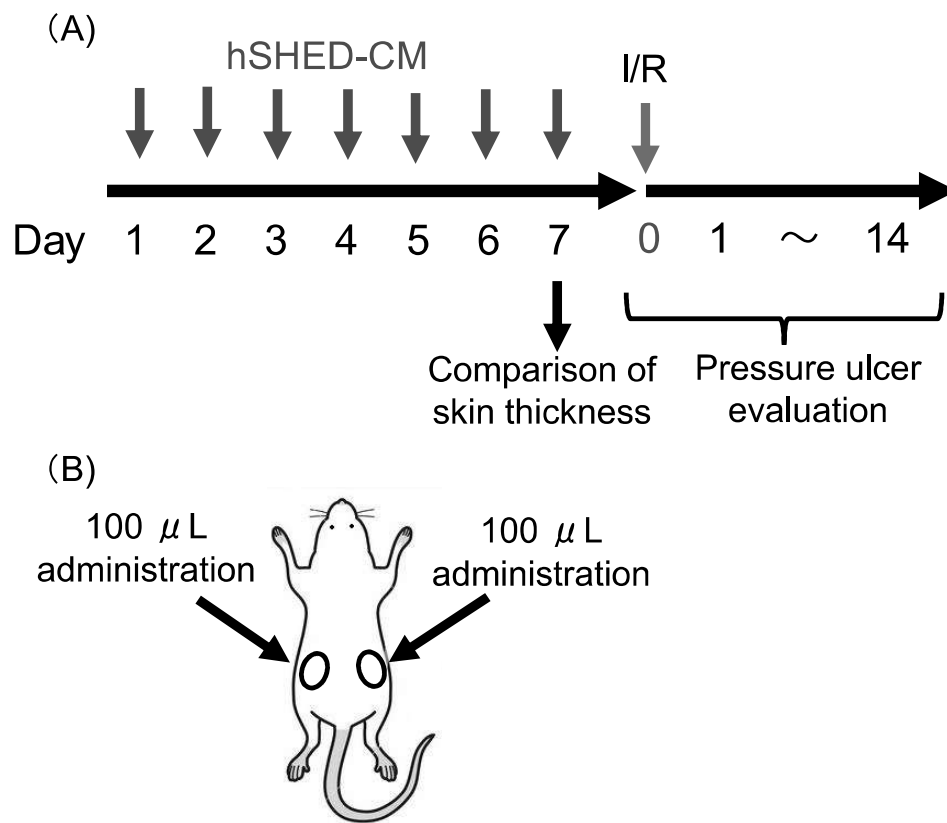


Fig. 27

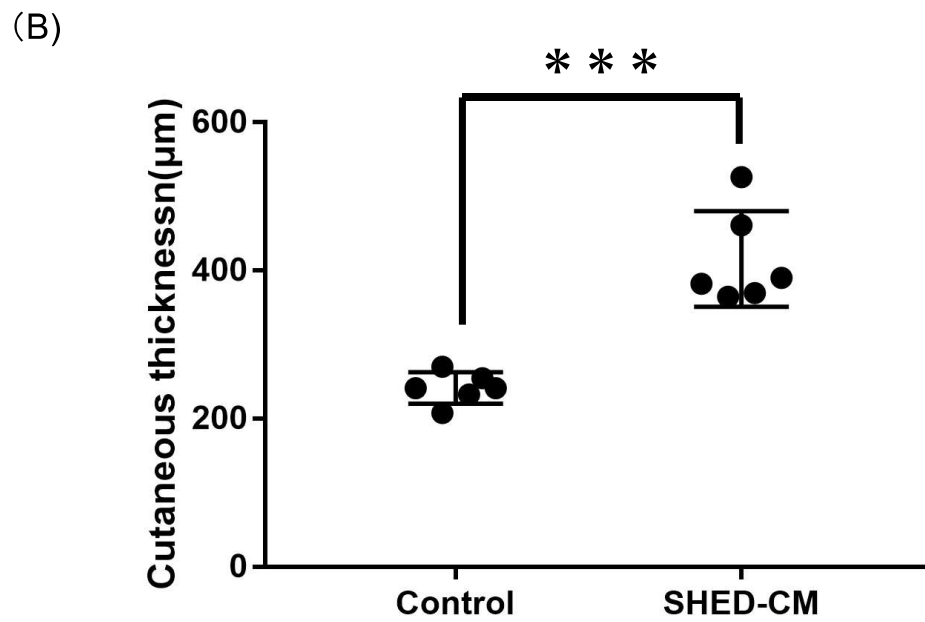
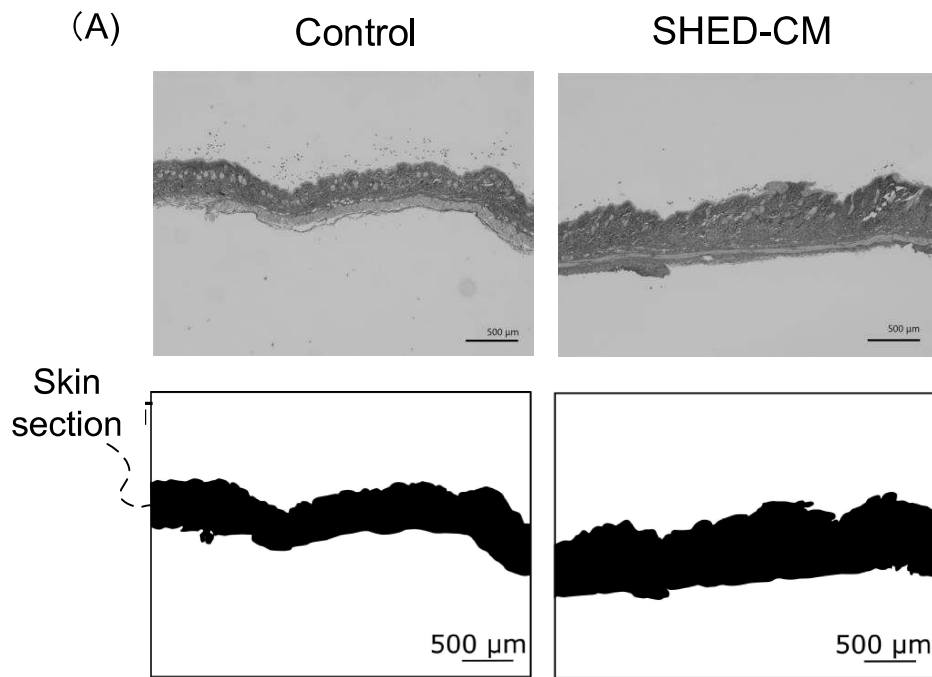
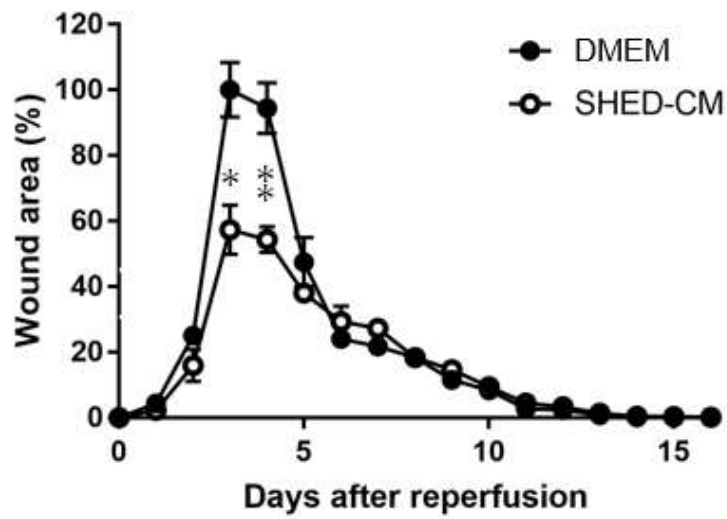


Fig. 28

(A)



(B)

