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CLUSTAL 2.1 multiple sequence alignment

Full            MASRWAVQLLLVAAWSMGCGEALKCYTCKEPMTSASCRTIIRCKPEDIACMTILVTVEAE 60
Mature          ----------------------LKCYTCKEPMTSASCRTIIRCKPEDIACMTILVTVEAE 38
                                       ************************************

Full            YPFNQSPVVTRSCSSSCVAIDPDSIGAAHLIFCCFRDLCNSEL 103  SEQ ID NO:2
Mature          YPFNQSPVVTRSCSSSCVAIDPDSIGAAHLIFCCFRDLCNSEL 81 SEQ ID NO:1
                ********************************************
```

FIG. 1

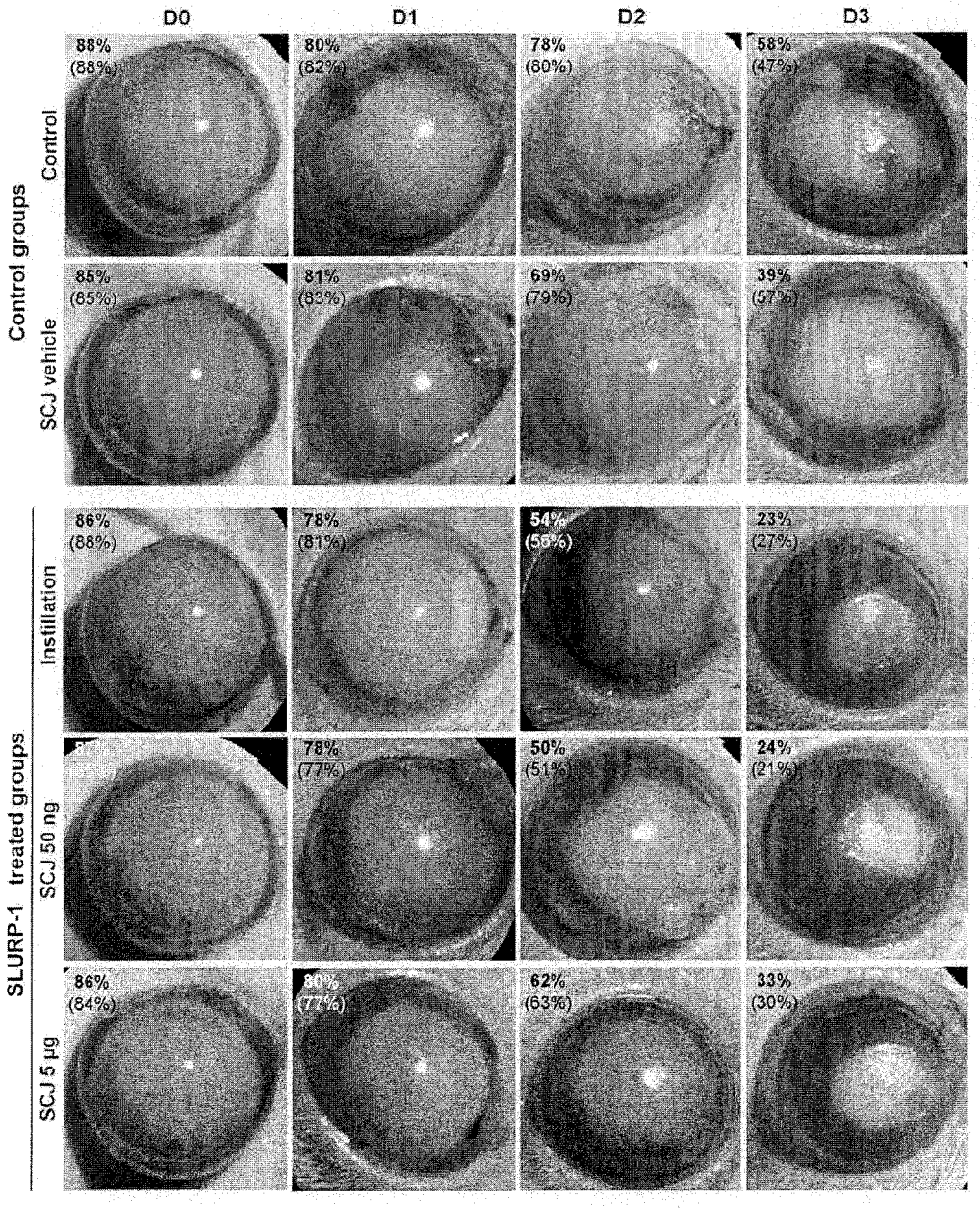



FIG. 2

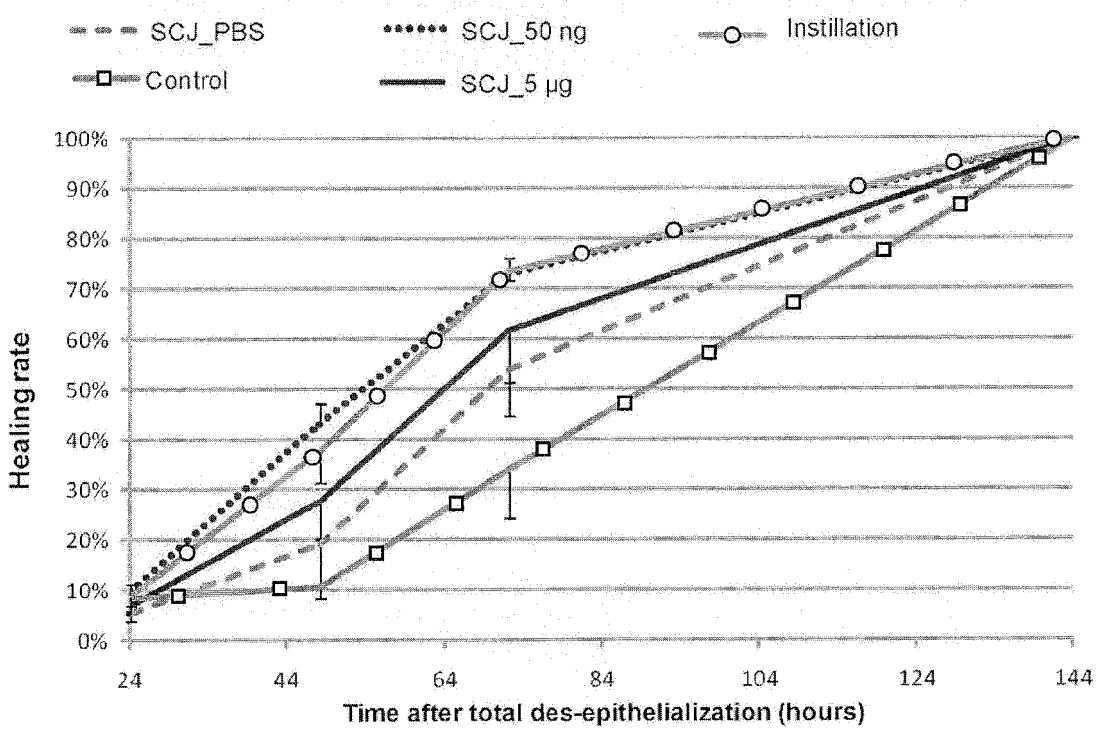



FIG. 3a

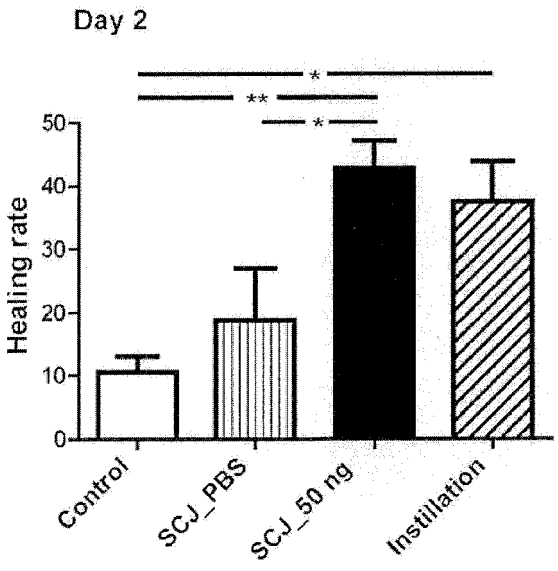



FIG. 3b

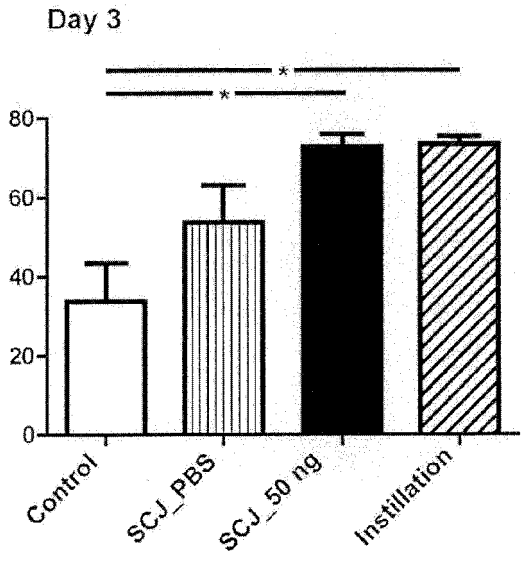



FIG. 3c

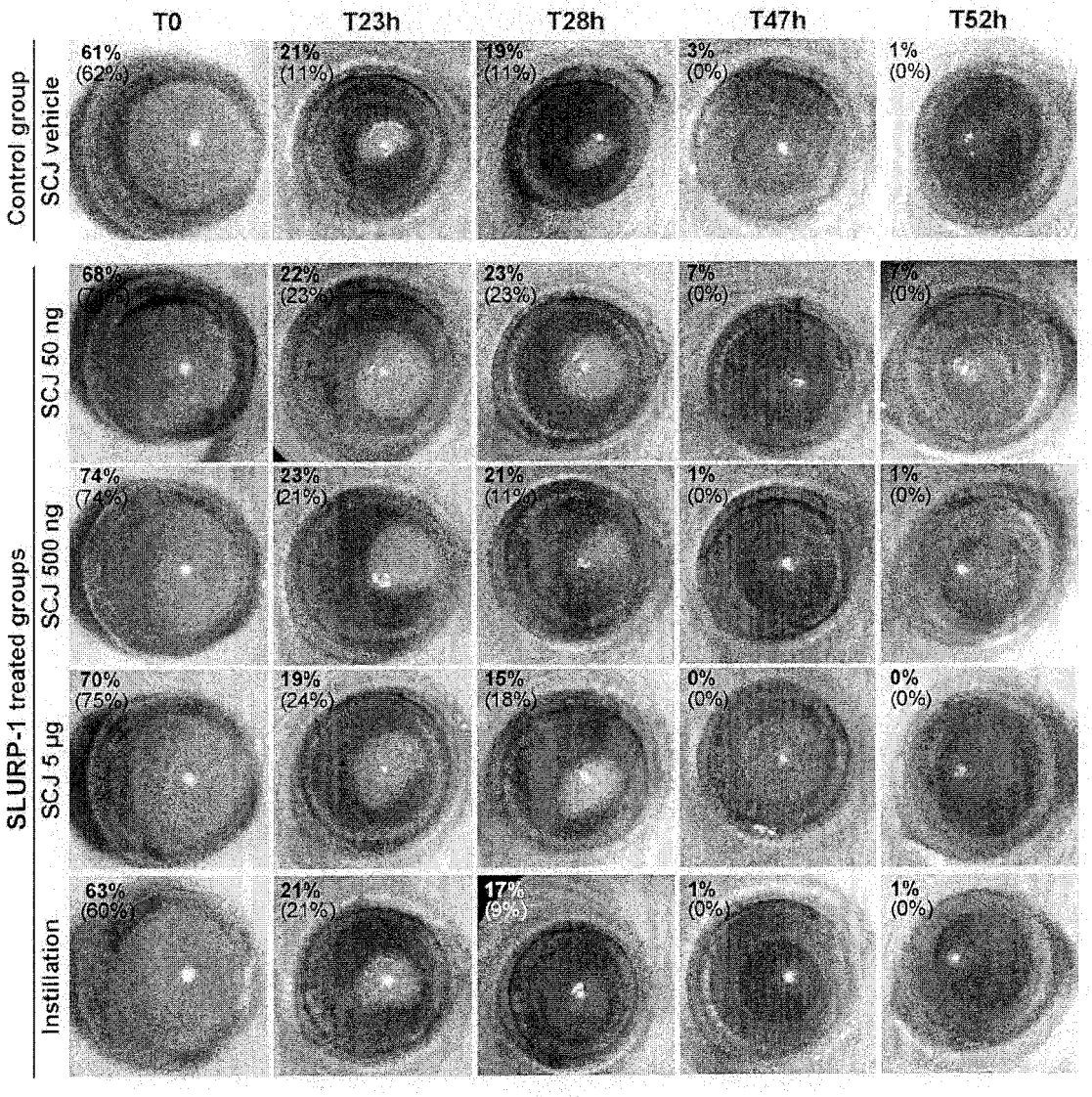

T0
T23h
T28h
T47h
T52h
Control group
SCJ vehicle
61% (62%)
21% (11%)
19% (11%)
3% (0%)
1% (0%)
SLURP-1 treated groups
SCJ 50 ng
68% (74%)
22% (23%)
23% (23%)
7% (0%)
1% (0%)
SCJ 500 ng
74% (74%)
23% (21%)
21% (11%)
1% (0%)
1% (0%)
SCJ 5 µg
70% (75%)
19% (24%)
15% (18%)
0% (0%)
0% (0%)
Instillation
63% (60%)
21% (21%)
17% (9%)
1% (0%)
1% (0%)

FIG. 4

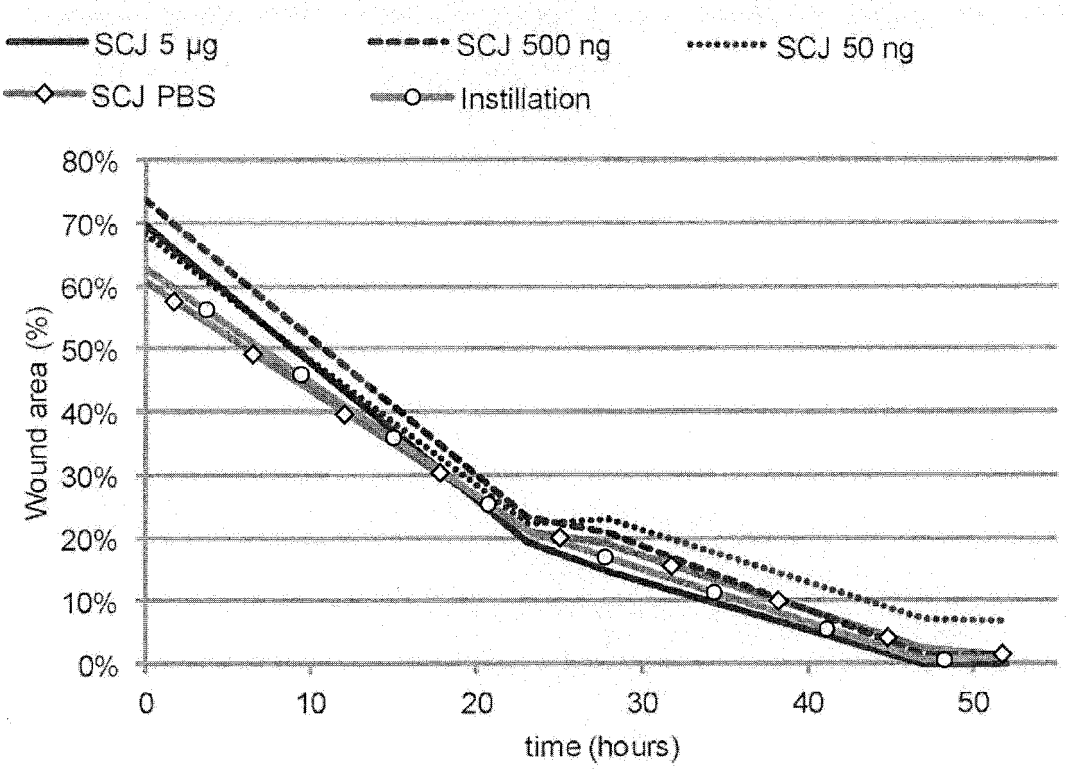



FIG. 5a

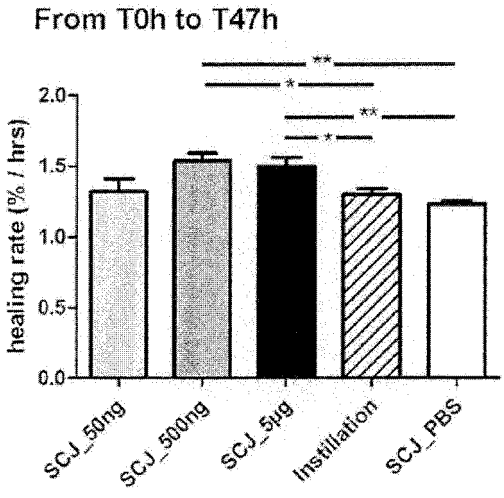



FIG. 5b

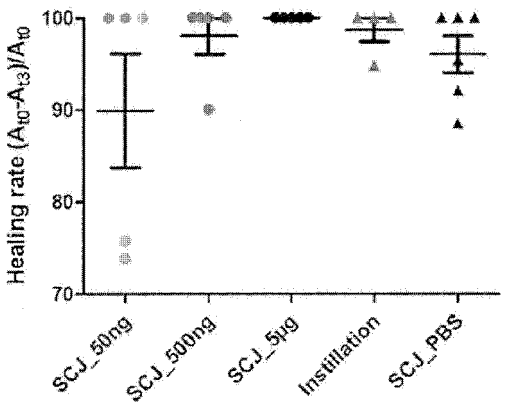
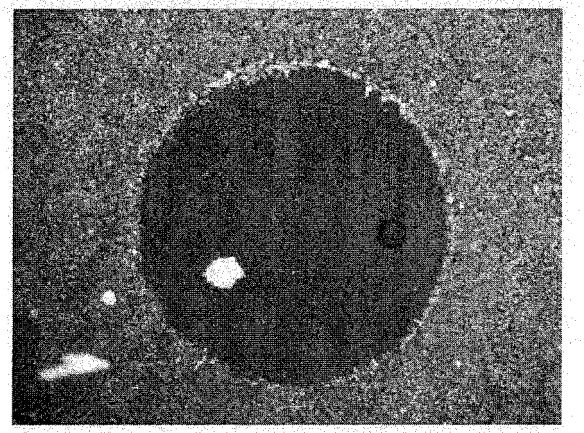
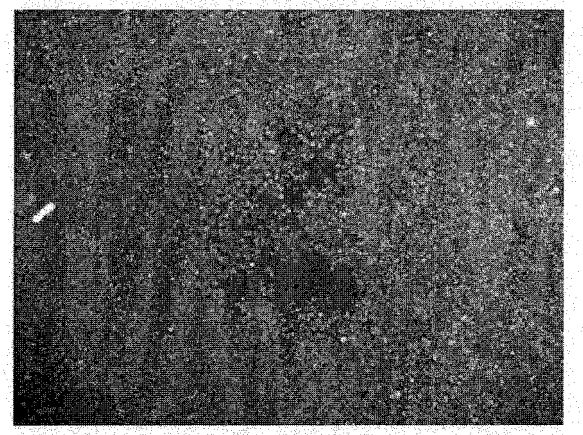



FIG. 5c

100% open wound, 0% wound closure



10 µg/ml SLURP-1



Control (No treatment)

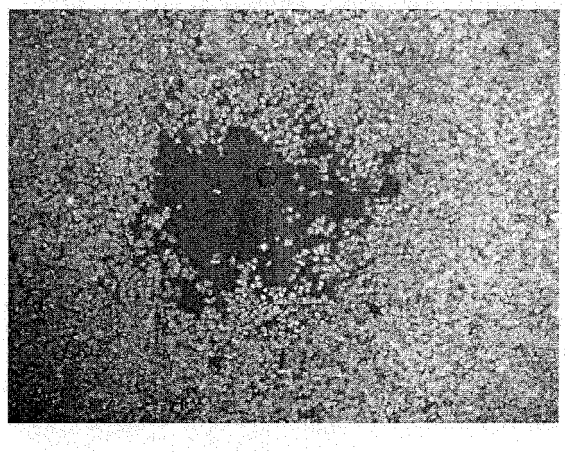


FIG. 6