

Article

Monocyte Chemotactic Protein-1 (MCP1) Accumulation in Human Osteoclast Precursor Cultures

Nigel A. Morrison ^{1,*} and Mark R. Forwood ²¹ School of Medical Science, Griffith University, Queensland, Australia; N.Morrison@griffith.edu.au² School of Medical Science, Griffith University, Queensland, Australia; M.Forwood@griffith.edu.au

* Correspondence: N.Morrison@griffith.edu.au

Supplementary Materials

Citation: Morrison, N.A.; Forwood, M.R. Monocyte Chemotactic Protein-1 (MCP1) Accumulation in Human Osteoclast Precursor Cultures. *Life* **2022**, *12*, 789. <https://doi.org/10.3390/life12060789>

Academic Editor: Christina Piperi

Received: 17 March 2022

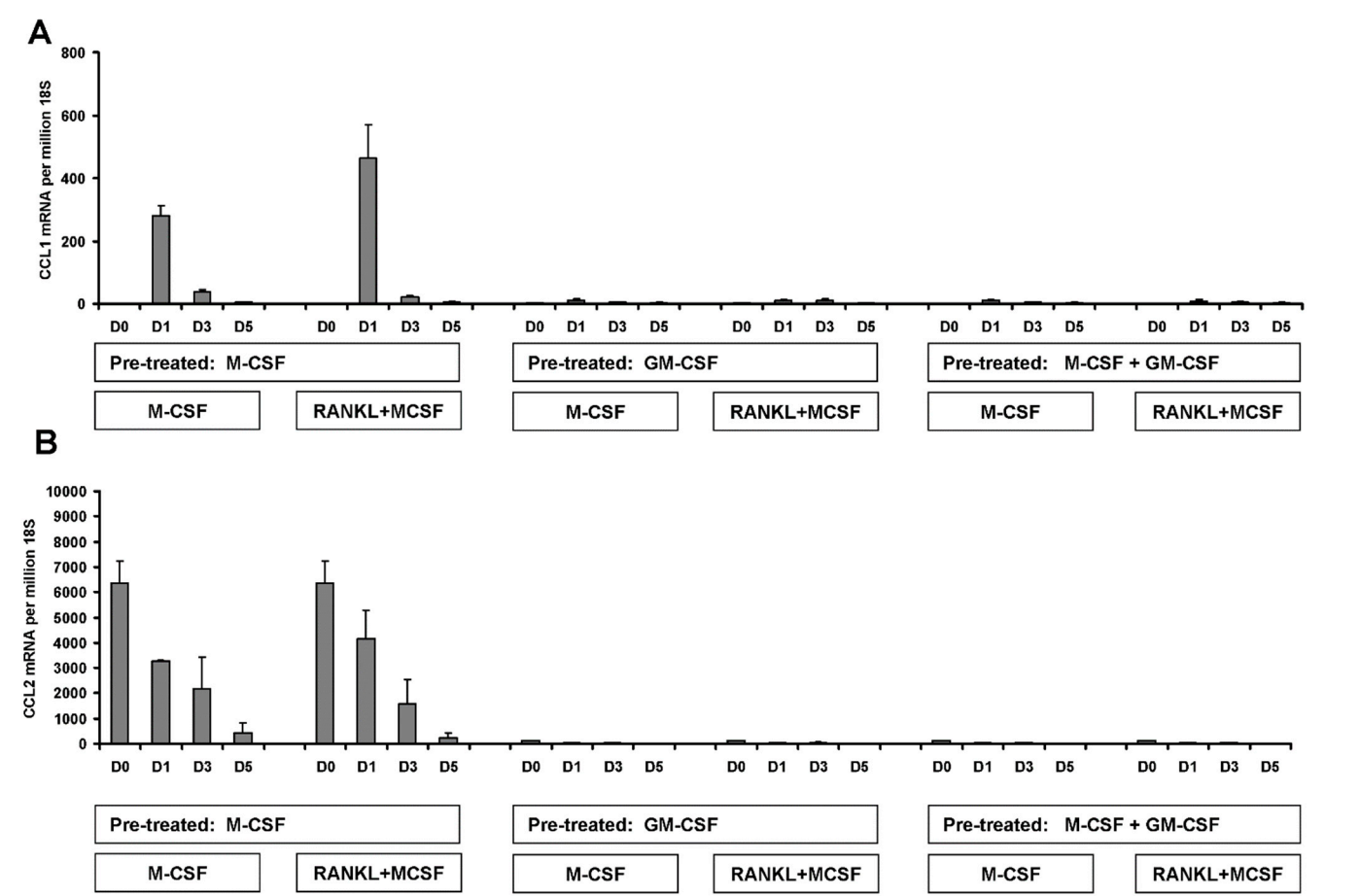
Accepted: 23 May 2022

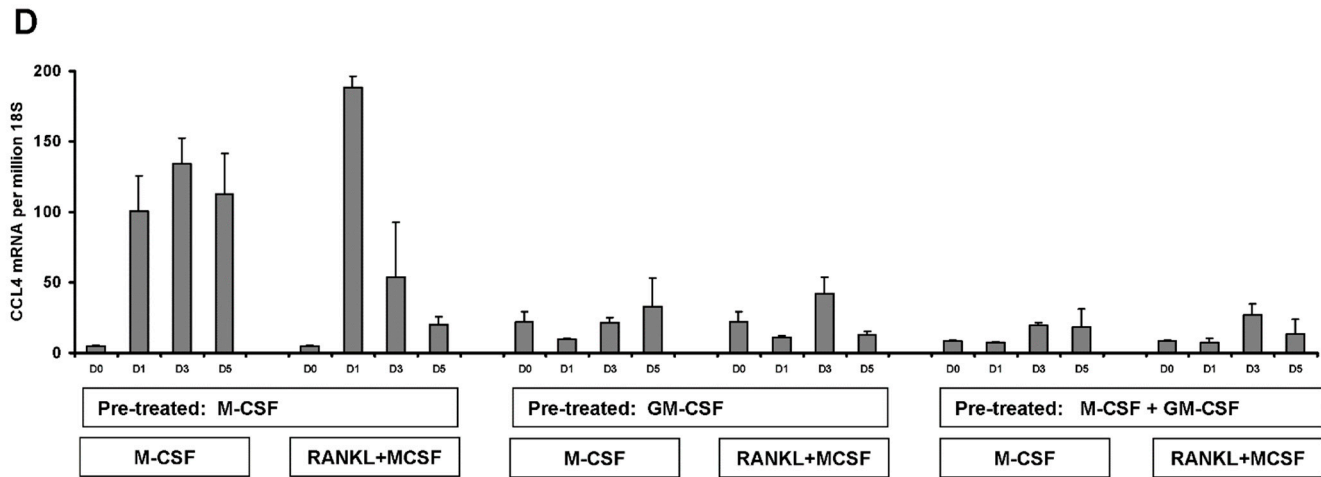
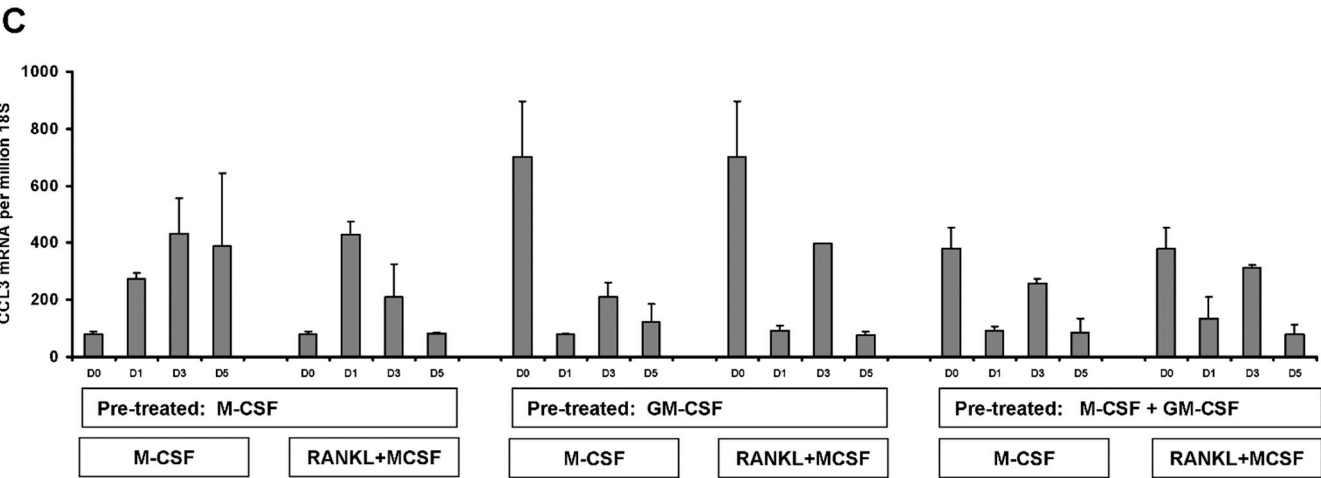
Published: 26 May 2022

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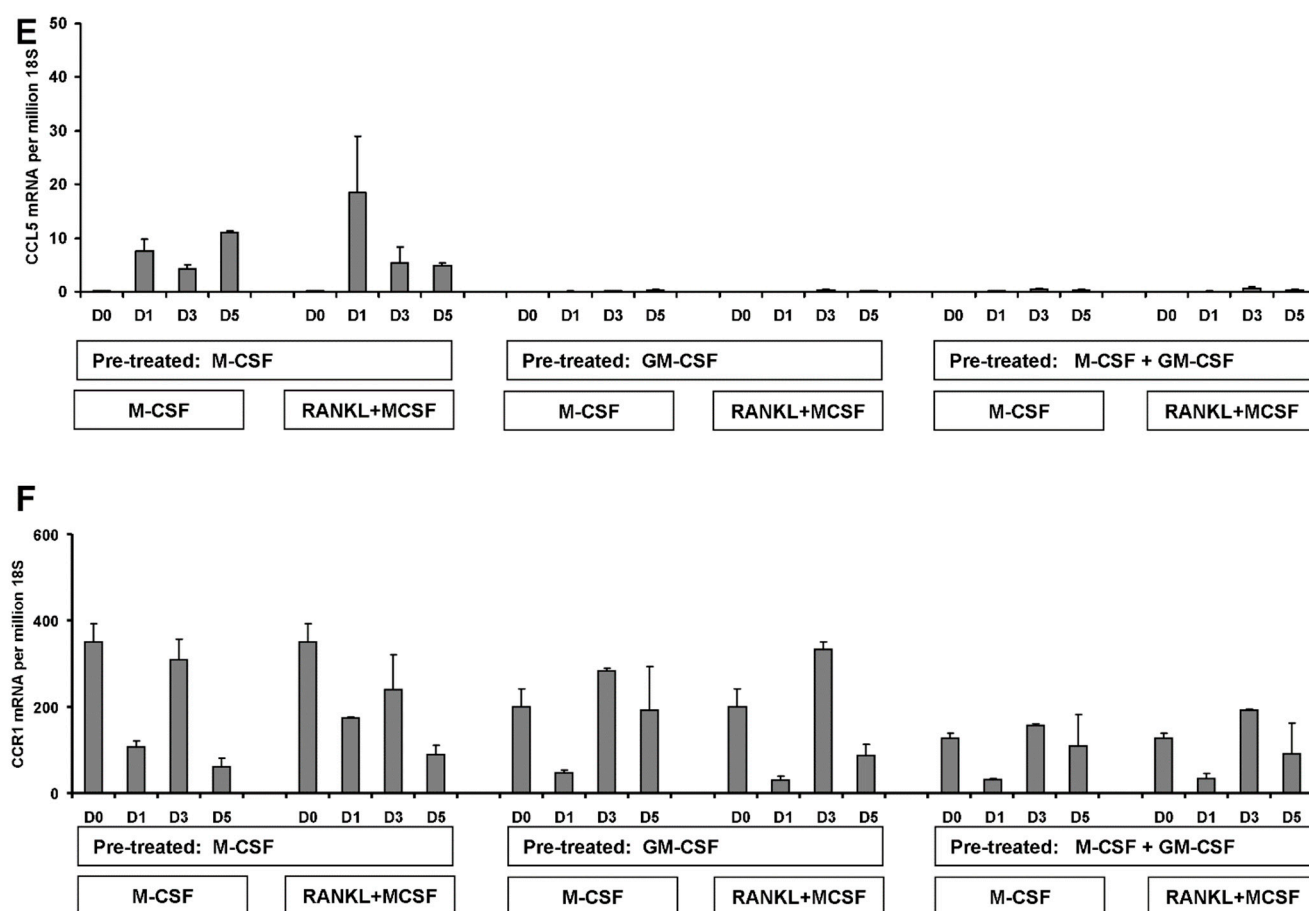


Figure S1. Chemokine mRNA transcript changes according to culture conditions. Each graph shows the outcome of six different conditions, with gene expression measured at zero time (D0), and then at days 1, 3 and 5 (D1, D3 and D5, respectively). Pre-treatment for 5 days is indicated in the first row of boxes containing the legend “Pre-treated”. Post treatment is indicated in the second set of boxes labelled either M-CSF or RANKL+M-CSF. **A.** CCL1 mRNA levels are very low except at day one after media change and only in those cultures pre-treated with M-CSF. **B.** MCP1 (CCL2) mRNA transcript levels are the highest of all assayed genes and are substantially suppressed by GM-CSF pre-treatment (zero time). High MCP1 mRNA levels in M-CSF pre-treated cells decline steadily over time. In GM-CSF pre-treated cells, no recovery of MCP1 mRNA levels was observed when cells were cultured in M-CSF alone or M-CSF in combination with RANKL. **C.** CCL3 (MIP1 α) mRNA levels are quite different at zero time (D0) according to pre-treatment with highest levels in GM-CSF treated cells. In cells pre-treated with M-CSF, an increase in expression is observed at days 1, 3 and 5 in both post-treatments of M-CSF and MCSF and RANKL. In GM-CSF pre-treated cells, CCL3 mRNA levels drop after day zero. **D.** CCL4 (MIP1 β) has very low transcript levels at day zero (immediately after the pre-treatment period). In M-CSF post-treatment cultures and M-CSF and RANKL post-treatment cultures, CCL4 (MIP1 β) shows induced gene expression effect in a pattern similar to that of CCL3 (MIP1 α). **E.** CCL5 (RANTES) mRNA levels are extremely low with highest levels in cultures pre-treated with M-CSF. **F.** CCR1 mRNA transcript levels are not particularly altered and appear to vary in all cultures.