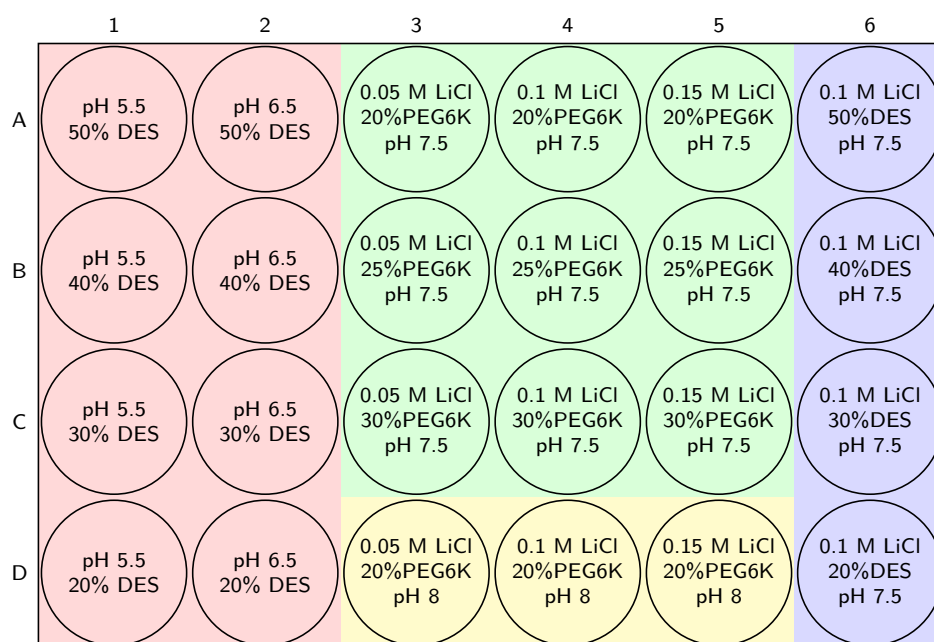


# WEB328 - PFOMT Xtallization

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**Figure 1:** The plate layout for crystallization of proteinase K, lipase B and PFOMT.

| component              | concentration         | volume       | stock                   |
|------------------------|-----------------------|--------------|-------------------------|
| SAE                    | 0.25 mM SAE           | 1 $\mu$ l    | 5 mM SAE                |
| MgCl <sub>2</sub>      | 0.25 mM               | 0.2 $\mu$ l  | 25 mM MgCl <sub>2</sub> |
| eriodictyol            | 0.25 mM               | 0.2 $\mu$ l  | 25 mM ED                |
| PFOMT                  | 0.262 mM (7.53 mg/mL) | 6.4 $\mu$ l  | 24 mg/mL PFOMT          |
| ad to 10 $\mu$ l water |                       | 12.2 $\mu$ l |                         |

**Table 1:** freshly prepared PFOMT xtallization solution.

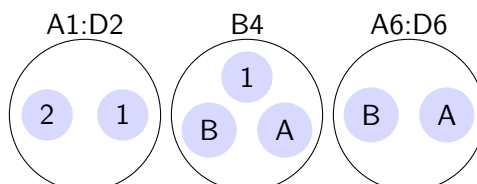
## 0.1 Buffers used:

A1:D1 → buffer is 0.1 M NaOAc

A2:D2 → buffer is 0.1 M Na-citrate

A3:D5 → buffer is 0.1 M Hepes

A6:D6 → buffer is 0.1 M Hepes



**Figure 2:** Layout of glass plates in special rows. View from the top. In wells A1 to D1, drop 2 contains proteinase K, drop 1 contains *C. cylindrica* lipase B. In wells A2 to D2, the order of drops is switched. Drop 1 contains Proteinase K and drop 2 contains lipase B. In well B5 drop 1 contains freshly prepared PFOMT (see WEB327) with ED and SAE. Drops B and A contain previously prepared PFOMT xtallization recipes (from WEB293). Wells A6:D6 also contain these preparations. A – 261  $\mu$ M SAH,  $\text{MgCl}_2$ , ferulic acid and 7.5 mg/mL PFOMT in 10 mM Tris pH 7.5. B – 0.5 mM SAE, 1 mM  $\text{MgCl}_2$ , 10 mg/mL PFOMT.

| Well  | Drop      | Protein      |
|-------|-----------|--------------|
| A1:D1 | 1 (right) | lipase B     |
|       | 2 (left)  | proteinase K |
| A2:D2 | 1 (right) | proteinase K |
|       | 2 (left)  | lipase B     |

**Table 2:** Protein in wells.

## 0.2 Method

500  $\mu$ l of crystallization buffer was used inside a well. On the glass slide used for crystallization, 1  $\mu$ l of protein solution was mixed with 1  $\mu$ l of buffer solution. The small end of a cat whisker was used for seeding. The whisker was dragged along the edges of a previous crystal and then through the crystallization drops. Afterwards the glass slide was put onto the well and placed at 4°C for crystallization.

## 0.3 Observations

**Day 0 - directly after seeding:** The drop 1 in wells B1 and C1 already contained crystals shortly after seeding. The drop 2 in wells D2 and C2 also already contained crystals along the seeding line.

**Day 4 after seeding:** In each well of row 1 and 2 the drops 1 and 2 contain crystals respectively.