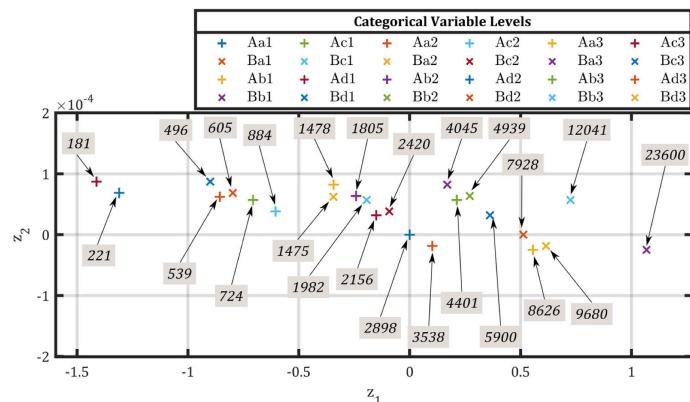


# DOE Meeting 05022024

Thursday, May 2, 2024 12:35 PM

LMGP combined with non-hierarchical multitask learning



<https://bpb-us-e2.wpmucdn.com/faculty.sites.uci.edu/dist/1/863/files/2023/01/LMGP-Paper.pdf>

$X = [X_t, X_d]$

$Z = A X_t$

$X' = [Z, X_d]$

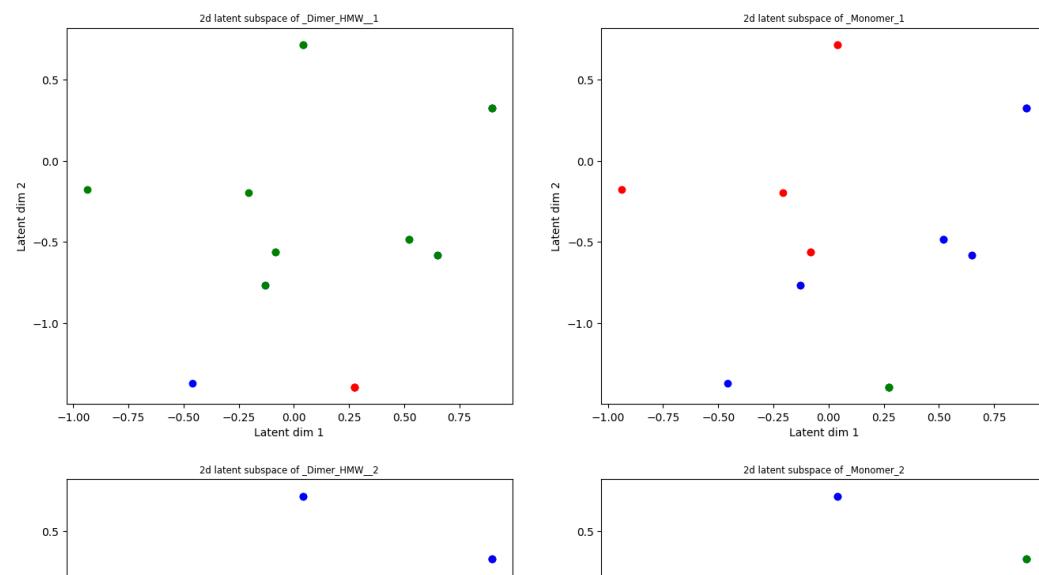
Input

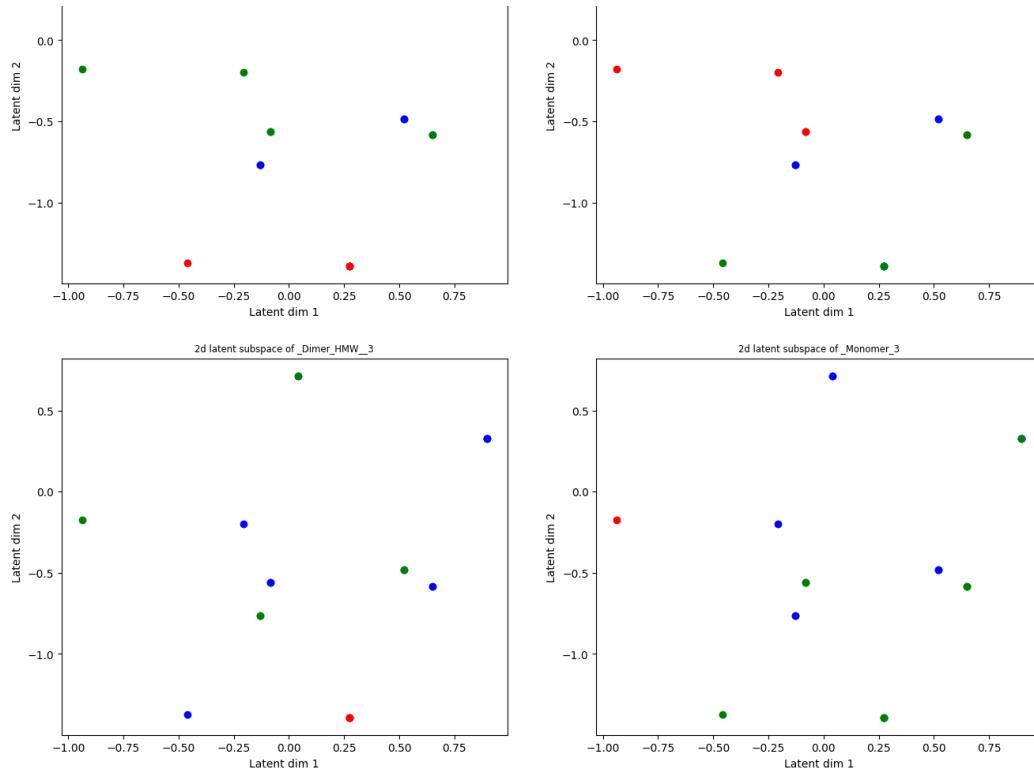
Formulation #	Buffer Type	pH	Sugar/ Salt	Additive	Additive Conc (mM)	PS20 Conc (%)
1	Ace	4.5	4% Sorbitol	Arginine	40	0.02
2	Ace	4.5	4% Sorbitol	Glycine	40	0.02
3	Ace	5	4% Sorbitol	Arginine	40	0.02
4	Ace	5	4% Sorbitol	Glycine	40	0.02
5	Ace	5	0.1M NaCl	Glycine	40	0.02
6	Ace	5.5	4% Sorbitol	Glycine	40	0.02
7	His	5.5	4% Sorbitol	Arginine	40	0.02
8	His	6	4% Sorbitol	Arginine	40	0.02
9	His	6	4% Sorbitol	Glycine	40	0.02
10	His	6	0.1M NaCl	Glycine	40	0.02
11	His	6.5	4% Sorbitol	Arginine	40	0.02
12	His	6.5	4% Sorbitol	Glycine	40	0.02
13	Cit	5	4% Sorbitol	Arginine	40	0.02
14	Cit	5	0.1M NaCl	Glycine	40	0.02
15	Cit	6	4% Sorbitol	Glycine	40	0.02
16	Cit	6.5	4% Sorbitol	Arginine	40	0.02
17	Cit	6.5	4% Sorbitol	Glycine	40	0.02
18	Cit	7	0.1M NaCl	Glycine	40	0.02
19	Cit	7	4% Sorbitol	Glycine	40	0.02
20	Cit	7	4% Sorbitol	Arginine	40	0.02
21	None	6.5	4% Sorbitol	Arginine	40	0.02

z1	Phos	0.5	4% Sorbitol	Arginine	40	0.02
22	Phos	6.5	4% Sorbitol	Glycine	40	0.02
23	Phos	7	4% Sorbitol	Arginine	40	0.02
24	Phos	7	4% Sorbitol	Glycine	40	0.02

### Output

	%Dimer (HMW)		%Monomer			%LMW			
			40°C	T0	FT	40°C	T0	FT	40°C
1	2.4	1.6	1.8	95.1	96.2	93.8	2.5	2.2	4.4
2	0.8	1.1	0.7	97.3	96.6	95.1	2.0	2.2	4.2
3	3.2	2.0	2.7	94.3	95.6	93.4	2.6	2.4	3.9
4	1.4	1.4	1.3	96.7	96.4	95.0	1.9	2.3	3.6
5	4.2	2.3	3.4	93.3	95.2	92.7	2.5	2.4	4.0
6	1.8	1.3	1.9	96.3	96.4	94.5	2.0	2.3	3.6
7	3.6	2.2	3.1	94.0	95.5	93.8	2.4	2.3	3.1
	<b>T0</b>								
	<b>FT</b>								
8	3.7	2.1	3.4	94.3	95.6	93.0	1.9	2.4	3.5
9	2.1	1.3	2.2	95.5	96.4	94.2	2.4	2.3	3.6
<b>10</b>	4.3	2.1	4.9	93.1	95.5	89.6	2.6	2.4	5.5
11	4.4	2.1	4.3	93.3	95.8	92.1	2.4	2.1	3.6
12	1.9	1.2	2.8	96.1	96.8	93.3	2.0	2.0	3.9
13	2.9	1.5	2.5	95.1	96.3	93.6	2.0	2.2	4.0
<b>14</b>	3.5	1.8	3.4	94.5	95.9	92.4	2.0	2.3	4.2
15	3.2	1.9	3.0	94.8	96.0	93.4	2.0	2.1	3.6
<b>16</b>	3.2	1.7	3.1	94.8	96.0	93.0	1.9	2.2	3.9
<b>17</b>	3.3	1.9	3.3	94.7	95.8	92.9	2.0	2.3	3.9
<b>18</b>	3.7	1.9	4.0	94.3	95.9	91.5	2.0	2.2	4.5
<b>19</b>	3.3	1.8	3.6	94.7	95.9	91.8	2.0	2.3	4.6
<b>20</b>	3.3	1.7	3.3	94.4	96.2	92.3	2.4	2.1	4.4
<b>21</b>	3.4	1.8	3.7	94.1	96.0	92.3	2.5	2.2	4.0
<b>22</b>	3.0	1.9	3.6	94.4	95.7	92.2	2.5	2.5	4.2
<b>23</b>	3.4	1.8	4.2	94.1	95.9	90.8	2.5	2.3	5.0
<b>24</b>	3.3	1.9	4.4	94.6	95.9	90.3	2.2	2.2	5.3





MAE	<u>Dimer_HMW_1</u>	<u>Monomer_1</u>	<u>Dimer_HMW_2</u>	<u>Monomer_2</u>	<u>Dimer_HMW_3</u>	<u>Monomer_3</u>
No LMGP	0.22	0.46	0.1	0.21	0.43	0.69
LMGP	0.49	0.54	0.18	0.28	<b>0.22</b>	<b>0.29</b>
Hier LMGP	0.19	0.21	0.06	0.16	0.31	0.54

Avg STD	<u>Dimer_HMW_1</u>	<u>Monomer_1</u>	<u>Dimer_HMW_2</u>	<u>Monomer_2</u>	<u>Dimer_HMW_3</u>	<u>Monomer_3</u>
No LMGP	1.1	1.27	0.49	0.61	1.34	2.06
LMGP	<u>0.37</u>	<u>0.49</u>	<u>0.18</u>	<u>0.26</u>	<u>0.41</u>	<u>0.66</u>
Hier LMGP	0.27	0.49	0.23	0.42	0.26	0.46

