

The dogs data

February 22, 2011

The data:

| | | | | | |
|--------------|---|-------|-------|-------|-------|
| Morphine | N | -3.22 | -1.61 | -2.30 | -2.53 |
| Morphine | N | -3.91 | -2.81 | -3.91 | -3.91 |
| Morphine | N | -2.66 | 0.34 | -0.73 | -1.43 |
| Morphine | N | -1.77 | -0.56 | -1.05 | -1.43 |
| Trimethaphan | N | -3.51 | -0.48 | -1.17 | -1.51 |
| Trimethaphan | N | -3.51 | 0.05 | -0.31 | -0.51 |
| Trimethaphan | N | -2.66 | -0.19 | 0.07 | -0.22 |
| Trimethaphan | N | -2.41 | 1.14 | 0.72 | 0.21 |

The SAS code and output:

```
options linesize=75;
```

```
data dogs;
  infile "dogs.dat";
  input Drug $ Depleted $ Histamine0 Histamine1
        Histamine3 Histamine5;
  LogHistamine0=log(Histamine0);
  LogHistamine1=log(Histamine1);
  LogHistamine3=log(Histamine3);
  LogHistamine5=log(Histamine5);
```

```
proc glm;
  class Drug Depleted;
  model LogHistamine0--LogHistamine5 =
        Drug Depleted Drug*Depleted / noint;
  repeated Time 4 (0 1 3 5);
```