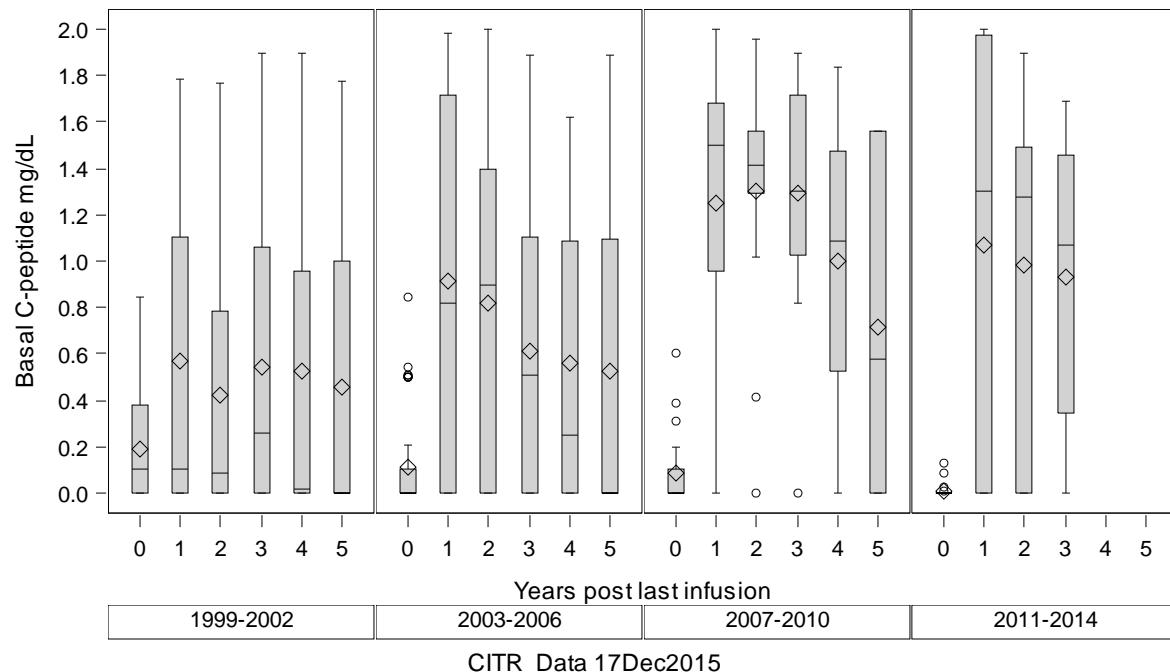


Exhibit A-10b. IAK Fasting C-peptide levels post last infusion (boxplots and generalized estimating equations)

Basal C-peptide (mg/dL) by Era



The GENMOD Procedure

Analysis Of GEE Parameter Estimates

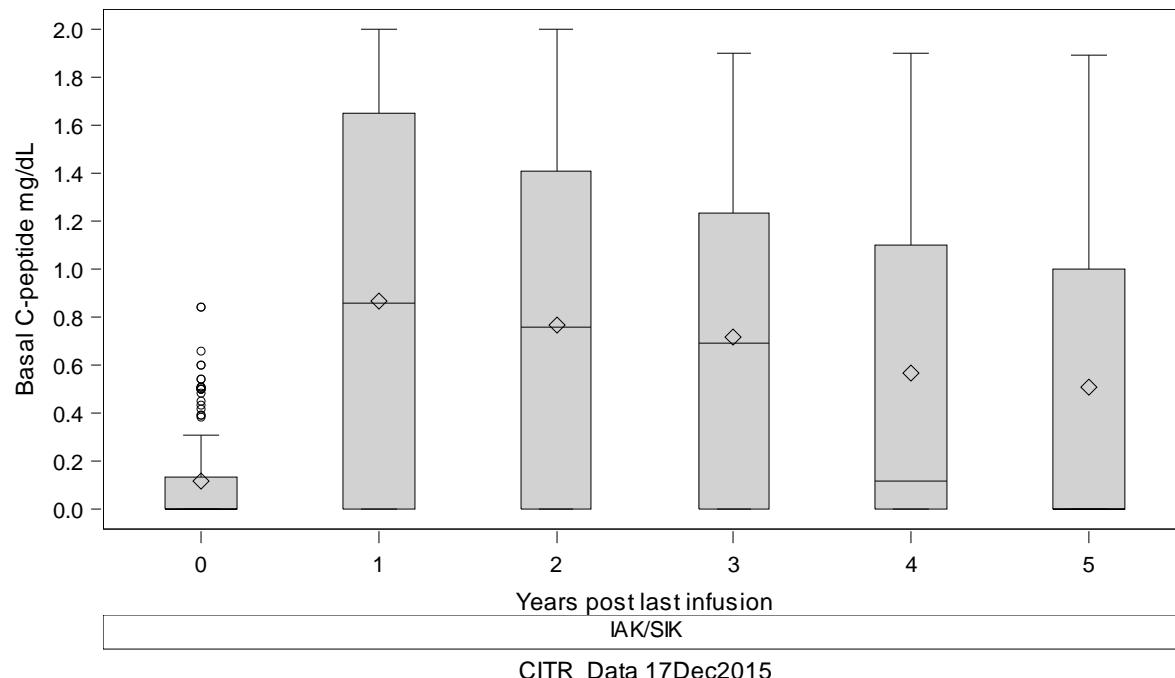
Empirical Standard Error Estimates

| Parameter | Estimate | Standard Error | 95% Confidence Limits | | Z | Pr > Z |
|-----------|----------|----------------|-----------------------|---------|-------|---------|
| Intercept | -0.6433 | 0.2149 | -1.0646 | -0.2221 | -2.99 | 0.0028 |
| era | 0.2942 | 0.0797 | 0.1379 | 0.4504 | 3.69 | 0.0002 |
| yrspli | 0.0025 | 0.0224 | -0.0413 | 0.0464 | 0.11 | 0.9092 |

| Era | 1 | | 2 | | 3 | | 4 | | 5 | | | |
|-----------|--------|----------|--------|----|----------|-----|--------|----------|--------|----|----------|-----|
| | bcpep_ | | bcpep_ | | bcpep_ | | bcpep_ | | bcpep_ | | | |
| | N | Mea n | SE | N | Mea n | SE | N | Mea n | SE | N | Mea n | SE |
| 1999-2002 | 42 | 1.3 | 0.2 | 39 | 1.1 | 0.2 | 38 | 0.8 | 0.2 | 33 | 0.7 | 0.2 |
| 2003-2006 | 48 | 1.5 | 0.2 | 44 | 1.3 | 0.2 | 40 | 1.0 | 0.2 | 32 | 0.8 | 0.2 |
| 2007-2010 | 25 | 2.3 | 0.3 | 22 | 2.4 | 0.3 | 21 | 2.6 | 0.5 | 8 | 2.0 | 0.5 |
| 2011-2014 | 18 | 2.1 | 0.5 | 15 | 1.7 | 0.4 | 10 | 1.3 | 0.3 | . | . | . |

Exhibit A-10b. IAK Fasting C-peptide levels post last infusion (boxplots and generalized estimating equations)

Basal C-peptide (mg/dL) by Transplant type



The GENMOD Procedure

Analysis Of GEE Parameter Estimates

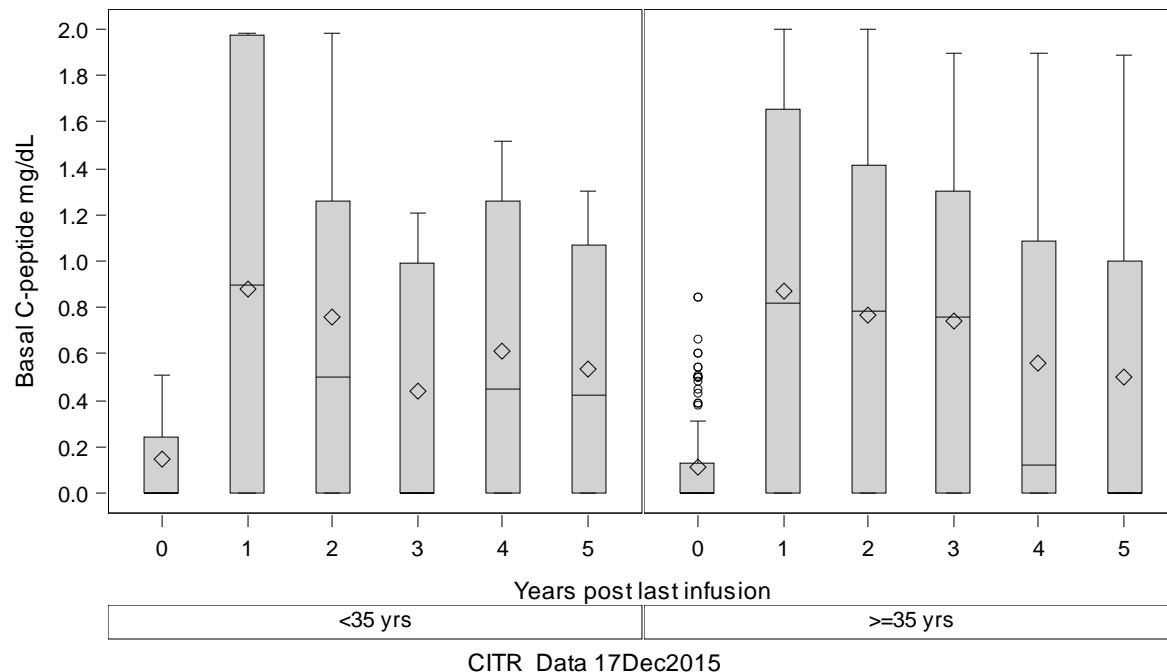
Empirical Standard Error Estimates

| Parameter | Estimate | Standard Error | 95% Confidence Limits | | Z | Pr > Z |
|------------|----------|----------------|-----------------------|--------|-------|---------|
| Intercept | 0.0847 | 0.0741 | -0.0605 | 0.2300 | 1.14 | 0.2530 |
| typeoftran | 0.0000 | 0.0000 | 0.0000 | 0.0000 | . | . |
| yrspli | -0.0387 | 0.0204 | -0.0788 | 0.0013 | -1.90 | 0.0580 |

| Transplant type | 1 | | | 2 | | | 3 | | | 4 | | | 5 | | |
|-----------------|--------|-----|-----|--------|-----|-----|--------|-----|-----|--------|-----|-----|--------|-----|-----|
| | bcpep_ | | | bcpep_ | | | bcpep_ | | | bcpep_ | | | bcpep_ | | |
| | Mea | n | SE |
| IAK | 133 | 1.7 | 0.1 | 120 | 1.5 | 0.1 | 109 | 1.3 | 0.1 | 73 | 0.9 | 0.1 | 65 | 0.9 | 0.2 |

Exhibit A-10b. IAK Fasting C-peptide levels post last infusion (boxplots and generalized estimating equations)

Basal C-peptide (mg/dL) by Recipient age



The GENMOD Procedure

Analysis Of GEE Parameter Estimates

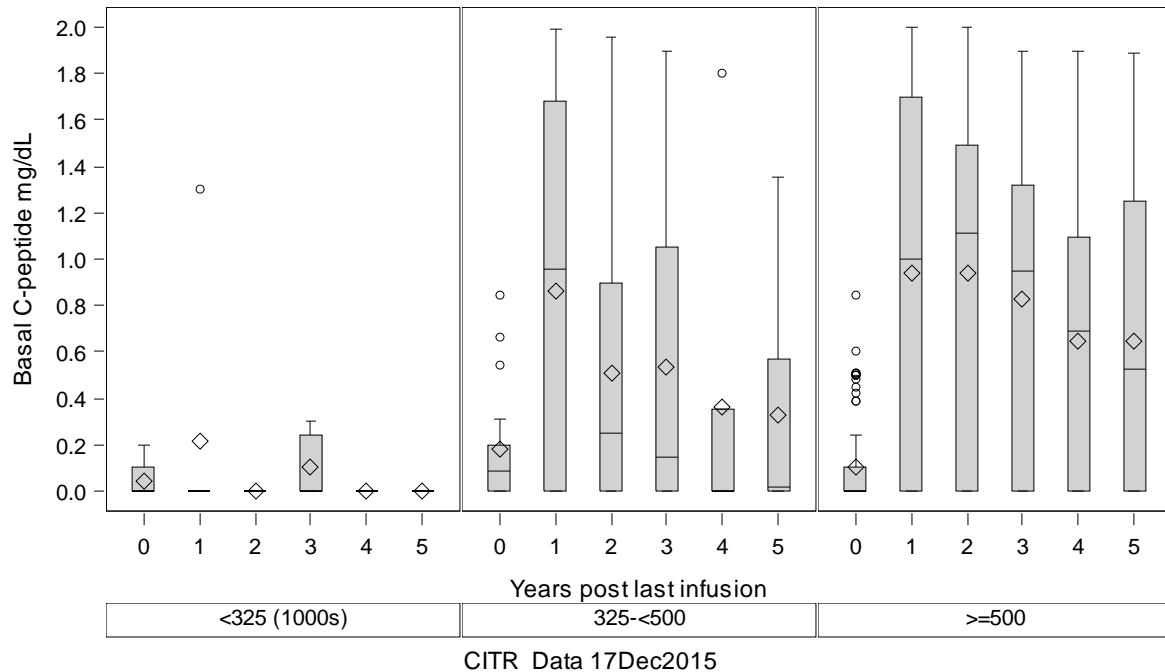
Empirical Standard Error Estimates

| Parameter | Estimate | Standard Error | 95% Confidence Limits | | Z | Pr > Z |
|-----------|----------|----------------|-----------------------|--------|-------|---------|
| Intercept | -0.2833 | 0.5447 | -1.3510 | 0.7843 | -0.52 | 0.6030 |
| age_c | 0.1913 | 0.2809 | -0.3592 | 0.7418 | 0.68 | 0.4959 |
| yrspli | -0.0380 | 0.0207 | -0.0786 | 0.0025 | -1.84 | 0.0659 |

| Recipient age | 1 | | 2 | | 3 | | 4 | | 5 | | | |
|---------------|--------|----------|--------|-----|----------|-----|--------|----------|--------|----|----------|-----|
| | bcpep_ | | bcpep_ | | bcpep_ | | bcpep_ | | bcpep_ | | | |
| | N | Mea n | SE | N | Mea n | SE | N | Mea n | SE | N | Mea n | SE |
| <35 yrs | 13 | 1.6 | 0.3 | 11 | 1.1 | 0.3 | 9 | 1.1 | 0.5 | 8 | 1.1 | 0.4 |
| ≥35 yrs | 120 | 1.7 | 0.1 | 109 | 1.5 | 0.1 | 100 | 1.3 | 0.2 | 65 | 0.9 | 0.1 |
| | | | | | | | | | | | | |
| | | | | | | | | | | | | |

Exhibit A-10b. IAK Fasting C-peptide levels post last infusion (boxplots and generalized estimating equations)

Basal C-peptide (mg/dL) by Total IEQs all infusions



The GENMOD Procedure

Analysis Of GEE Parameter Estimates

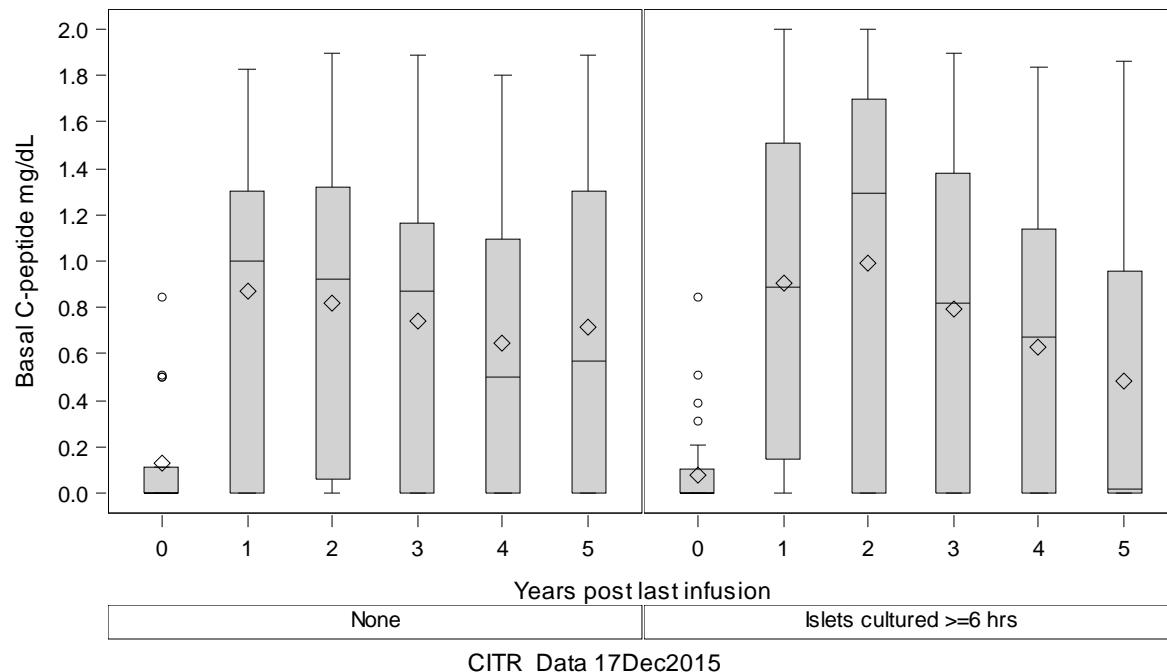
Empirical Standard Error Estimates

| Parameter | Estimate | Standard Error | 95% Confidence Limits | | Z | Pr > Z |
|-----------|----------|----------------|-----------------------|--------|-------|---------|
| Intercept | -1.4086 | 0.7886 | -2.9543 | 0.1371 | -1.79 | 0.0741 |
| ieqinf_c | 0.5445 | 0.2692 | 0.0169 | 1.0721 | 2.02 | 0.0431 |
| yrspli | -0.0364 | 0.0218 | -0.0791 | 0.0063 | -1.67 | 0.0943 |

| | 1 | | 2 | | 3 | | 4 | | 5 | | | |
|--------------------------|----------------|----------|----------------|----|----------------|-----|----------------|----------|----------------|----|----------|-----|
| | <i>bcppep_</i> | | <i>bcppep_</i> | | <i>bcppep_</i> | | <i>bcppep_</i> | | <i>bcppep_</i> | | | |
| | N | Mea n | SE | N | Mea n | SE | N | Mea n | SE | N | Mea n | SE |
| Total IEQs all infusions | | | | | | | | | | | | |
| <325 (1000s) | 7 | 0.6 | 0.4 | 6 | 1.0 | 1.0 | 6 | 1.0 | 0.9 | 4 | 0.0 | 0.0 |
| 325-<500 | 17 | 1.2 | 0.3 | 14 | 0.8 | 0.2 | 14 | 1.0 | 0.4 | 7 | 0.6 | 0.4 |
| >=500 | 90 | 1.9 | 0.2 | 84 | 1.7 | 0.2 | 73 | 1.5 | 0.2 | 50 | 1.0 | 0.2 |

Exhibit A-10b. IAK Fasting C-peptide levels post last infusion (boxplots and generalized estimating equations)

Basal C-peptide (mg/dL) by Cultured>6h



CITR Data 17Dec2015

The GENMOD Procedure

Analysis Of GEE Parameter Estimates

Empirical Standard Error Estimates

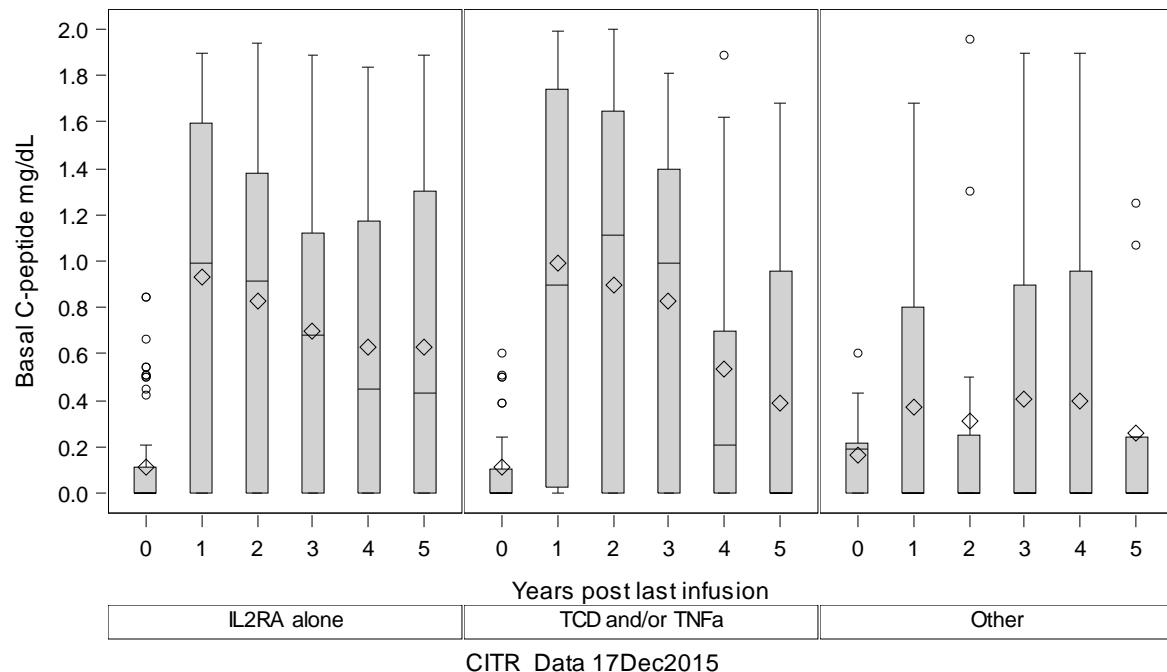
| Parameter | Estimate | Standard Error | 95% Confidence Limits | | Z | Pr > Z |
|-----------|----------|----------------|-----------------------|--------|-------|---------|
| Intercept | 0.0294 | 0.2048 | -0.3721 | 0.4308 | 0.14 | 0.8860 |
| cultured_ | 0.0388 | 0.2522 | -0.4555 | 0.5332 | 0.15 | 0.8777 |
| yrspli | -0.0311 | 0.0274 | -0.0847 | 0.0225 | -1.14 | 0.2561 |

| | 1 | | 2 | | 3 | | 4 | | 5 | | | |
|------------------------------------|---------------|----------|---------------|----|---------------|-----|---------------|----------|---------------|----|----------|-----|
| | <i>bcpеп_</i> | | <i>bcpеп_</i> | | <i>bcpеп_</i> | | <i>bcpеп_</i> | | <i>bcpеп_</i> | | | |
| | N | Mea n | SE | N | Mea n | SE | N | Mea n | SE | N | Mea n | SE |
| Cultured>6h | | | | | | | | | | | | |
| None | 19 | 1.6 | 0.3 | 18 | 1.0 | 0.2 | 17 | 1.2 | 0.3 | 15 | 0.9 | 0.2 |
| Islets cultured >= 6 hrs | 46 | 1.6 | 0.2 | 40 | 1.7 | 0.2 | 36 | 1.5 | 0.3 | 23 | 0.8 | 0.2 |

CITR Data 17Dec2015

Exhibit A-10b. IAK Fasting C-peptide levels post last infusion (boxplots and generalized estimating equations)

Basal C-peptide (mg/dL) by Induction



CITR Data 17Dec2015

The GENMOD Procedure

Analysis Of GEE Parameter Estimates

Empirical Standard Error Estimates

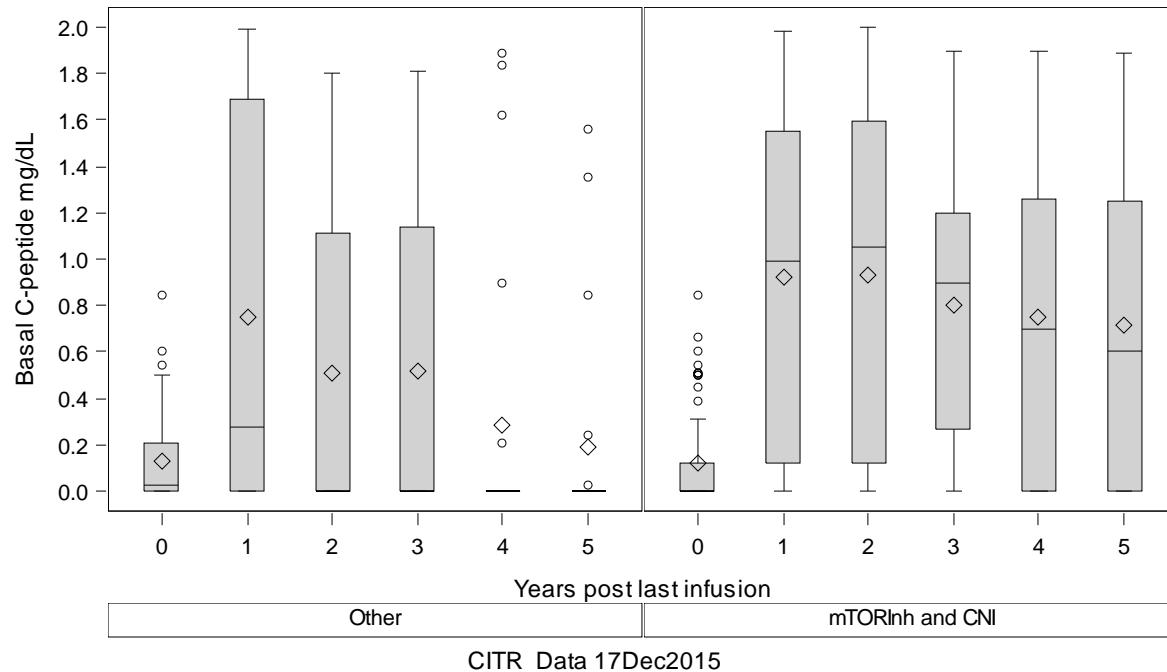
| Parameter | Estimate | Standard Error | 95% Confidence Limits | Z | Pr > Z |
|-----------|----------|----------------|-----------------------|-------|---------|
| Intercept | 0.1266 | 0.0924 | -0.0545 0.3077 | 1.37 | 0.1707 |
| inducco | -0.0518 | 0.1409 | -0.3279 0.2244 | -0.37 | 0.7133 |
| yrspli | -0.0389 | 0.0204 | -0.0789 0.0011 | -1.91 | 0.0565 |

| Induction | 1 | | 2 | | 3 | | 4 | | 5 | | | |
|-----------------|--------|------|--------|----|--------|-----|--------|------|--------|----|--------|-----|
| | bcpep_ | | bcpep_ | | bcpep_ | | bcpep_ | | bcpep_ | | bcpep_ | |
| | N | Mean | SE | N | Mean | SE | N | Mean | SE | N | Mean | SE |
| IL2RA alone | 62 | 1.6 | 0.2 | 61 | 1.2 | 0.2 | 57 | 1.3 | 0.2 | 42 | 0.9 | 0.2 |
| TCD and/or TNFa | 38 | 1.8 | 0.2 | 33 | 1.7 | 0.2 | 30 | 1.3 | 0.2 | 16 | 0.9 | 0.2 |
| Other | 19 | 1.6 | 0.5 | 17 | 1.7 | 0.6 | 16 | 1.3 | 0.5 | 13 | 0.9 | 0.4 |

CITR Data 17Dec2015

Exhibit A-10b. IAK Fasting C-peptide levels post last infusion (boxplots and generalized estimating equations)

Basal C-peptide (mg/dL) by Maintenance



The GENMOD Procedure

Analysis Of GEE Parameter Estimates

Empirical Standard Error Estimates

| Parameter | Estimate | Standard Error | 95% Confidence Limits | | Z | Pr > Z |
|-----------|----------|----------------|-----------------------|--------|-------|---------|
| Intercept | -0.2702 | 0.2009 | -0.6639 | 0.1235 | -1.34 | 0.1786 |
| maint | 0.5060 | 0.2206 | 0.0737 | 0.9384 | 2.29 | 0.0218 |
| yrspli | -0.0297 | 0.0212 | -0.0713 | 0.0118 | -1.40 | 0.1607 |

| Maintenance | 1 | | 2 | | 3 | | 4 | | 5 | | | |
|-----------------|--------|----------|--------|----|----------|-----|--------|----------|--------|----|----------|-----|
| | bcpep_ | | bcpep_ | | bcpep_ | | bcpep_ | | bcpep_ | | | |
| | N | Mea n | SE | N | Mea n | SE | N | Mea n | SE | N | Mea n | SE |
| Other | 46 | 1.5 | 0.2 | 42 | 1.2 | 0.2 | 39 | 0.9 | 0.2 | 25 | 0.4 | 0.2 |
| mTORinh and CNI | 73 | 1.8 | 0.2 | 69 | 1.6 | 0.2 | 64 | 1.5 | 0.2 | 46 | 1.2 | 0.2 |
| | | | | | | | | | | | | |